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RESULT 83
ACC45158/c
ID ACC451
XX ACC451
XX ACC451
XX Human;
XX Homo s
OS Synthe
XX Homo s
Synthe
YT modifi
FT modifi
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Best Local Similarity
Matches 20; Conserv
                                                                                                               New antisense compounds, useful for modulating the expression of NAC or for treating a disease or condition associated with the expression of NAC, e.g. hyperproliferative disease or neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; cytostatic; nootropic; neuroprotective; antiinflammatory; antisense therapy. NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. NAC is alsown as a death effector filament-forming CED4-like apoptosis protein (DEFCAR). NAC is located on human chromosome 17p13. The present sequence represents a human NAC chimeric phosphorothicate antisense oligonucleotide, which is given in the
                                                                                                                                                                                                                                                                                                   Bennett
                                                                                                                                                                                                                                                                                                                                                                                                               19-SEP-2001; 2001US-00956712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-SEP-2002; 2002WO-US029664.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO2003024988-A1.
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                                                            Exampel 15; Page 75; 147pp; English.
                                                                                                                                                                                                                                    WPI; 2003-354583/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
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                                                                                                                                                                                                                                                                                             Freier SM;
                                                                                                                                                                                                                                                                                                                                                      PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /*tag= b
/mod_base= OTHER
/mote= "2'-O-methoxyethyl (2'-MOB) gapmer"
/note= "2'-O-methoxyethyl (2'-MOB) gapmer"
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1. .5
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100.0%; Pr/
0;
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e= "2'-O-methoxyethyl (2'-MOB) gapmer"
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Pred. No.
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1.9e+02;
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The present invention describes a compound (I) 8-50 nucleobases in length

19-SEP-2002; 2002WO-US029664.

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ACC45170/c
ID ACC45170 standard; D
XX
AC ACC45170;
XC ACC45170;
XX
DT 16-JUN-2003 (first
ACC45170;
XX
DT 19-SEP-2002; 2002WO-
XX
DT 19-SEP-2002; 2002WO-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC targeted to a nucleic acid molecule encoding NAC, where the compound conhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid conlecule encoding NAC. Also described: (1) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC NAC in cells or tissues comprising contacting the cells or tissues with CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC NAC is inhibited. (I) has cytostatic, nootropic, neuroprotective and CC antisense compounds (I) are useful for modulating the expression of NAC, and for treating a disease or condition associated with expression of CC was prophylaxis, e.g. to prevent or delay infection, inflammation or CC useful as research reagents and kits, or for diagnostics, therapeutics CC and prophylaxis, e.g. to prevent or delay infection, inflammation or CC CED4-like apoptosis protein (DEFCAP). NAC is also known as a death effector filament-forming CC phosphorothioate antisense oligonucleotide, which is given in the cc exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; cytostatic; nootropic; neuroprotective; antiinflammatory; antisense therapy; NAC; DBFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 3 A; 7 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            phosphorothicate; ss.
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                                                                                                                                                                                            /note= '
                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                        /*tag= c
/mod_base= OTHER
                                                                                                                                                                                                                                                                                          note=
                                                                                                                                                                                                                                 mod_base=
                                                                                                                                                                                                                                                  *tag=
                                                                                                                                                                                                                                                                                                           mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.4%; Score 20;
100.0%; Pred. No.
                                                                                                                                                                                                               n3:-O-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                       "2'-O-methoxyethyl (2'-MOE) gapmer"
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                                                                                                                                                                                                                                                                                      phosphorothioate backbone"
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RESULT 85
ACC45172/c
ID ACC451
XX ACC451
XX ACC451
XX Human
DT 16-JUN
XX Human
XX Human;
KW Human;
KW death
KW neurol
KW phosph
XX Homo ©
OS Homo ©
OS Synthe
XX Synthe
XX Key
FT modifi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound CC specifically hybridises with the nucleic acid molecule encoding NAC and CC inhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid CC molecule encoding NAC. Also described: (1) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC (I); and (3) treating an animal having a disease or condition associated CC with NAC computsing administering (I) to the animal so that expression of CC (I); and (3) treating an animal having a disease or condition associated CC with NAC computsing administering (I) to the animal so that expression of CC antisnflammatory activities, and can be used in antisense therapy. The CC antisense compounds (I) are useful for modulating the expression of CC (NAC, e.g. hyperproliferative disease, neurological disease, or a disease CC or disorder arising from abberrant appotosis. The compounds are also useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or CC tumour formation. NAC is also known as a death effector filament-forming CC Expandition approach (DEFCAP). NAC is located on human chromosome CC phosphorothicate antisense oligonucleotide, which is given in the CC examplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                           antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                            ACC45172 standard;
                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                      Human; cytostatic; nootropic; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                 Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:32.
                                                                                                                                                                                                                                                                                                                                                          16-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                   ACC45172;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 5 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 75; 147pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAC, e.g. hyperproliferative disease or neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New antisense compounds, useful for modulating the expression of NAC for treating a disease or condition associated with the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-354583/33
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                                                              modified_base
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                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                  Location/Qualifiers
/note= "phosphorothioate backbone"
                   /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                            20
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RESULT 86
ACC45179/c
ID ACC451
XX
AC ACC451
XX
DT 16-JUN
XX
DE Human

ACC45179 standard; DNA; 20

ΒP

<u>,,</u>

16-JUN-2003 ACC45179

(first entry)

Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:39

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                                                                                                                                                                   The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound CC specifically hybridises with the nucleic acid molecule encoding NAC and Ci inhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid CC molecule encoding NAC. Also described: (I) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC (I); and (I) thas cytostatic, nootropic, neuroprotective and CC antiinflammatory activities, and can be used in antisense therapy. The CC antisense compounds (I) are useful for modulating the expression of NAC, C. e.g. hyperproliferative disease, neurological disease, or a disease or disorder arising from aberrant apoptosis. The compounds are also CC useful as research reagents and kits, or for diagnostics, therapeutics CC and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. NAC is also known as a death effector filament-forming CC CED4-like apoptosis protein (DEFCAP). NAC is located on human chromosome CC phosphorothioate antisense oligonucleotide, which is given in the ccemplification of the present invention
                                                                   Query Match
Best Local S
Matches 20
                                                                                                                                         Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New antisense compounds, useful for modulating the expression of NAC or for treating a disease or condition associated with the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-MAR-2003.
                                   2679 CCACTGCTTGTACGAGACTC 2698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     e.g. hyperproliferative disease or neurological disease.
20
                                                                                      Similarity
CCACTGCTTGTACGAGACTC 1
                                                                                                                                           BP; 5 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Freier
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/*tag= b

/mod_base= OTHER

/note= "2'-O-methoxyethyl (2'-MOE) gapmer"
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                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    _base≃ OTHER
                                                                                                           0.4%;
                                                                                      Score 20; Pred. No.
                                                                     0; Mismatches
                                                                                                           DB 1;
                                                                                      1.9e+02
                                                                                                         Length 20
                                                                         Indels
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The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound compound the proficially hybridises with the nucleic acid molecule encoding NAC and compound specifically hybridises with the compound specifically hybridises with the teast an 8-nucleobase portion of an active site on a nucleic acid compound specifically hybridises with the compound specifically hybridises with the compound specifically hybridises with the compound to the expression of the specific acid compound to the expression of the compound that the composition comprising (I) and (3) treating an animal having a disease or condition associated the NAC comprising administering (I) to the animal so that expression of the compounds (I) has cytostatic, notropic, neuroprotective and the specific antisense compounds (I) are useful for modulating the expression of NAC, and for treating a disease or condition associated with expression of the compounds (I) are useful for modulating the expression of NAC, and for treating a disease or condition associated with expression of the compounds are also conseill as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflament-forming CED4-like apoptosis protein (DEPCAP). NAC is located on human chromosome CC 17pl3. The present sequence represents a human NAC chimeric CC phosphorochioate antisense oligonucleotide, which is given in the compolification of the present invention
     Query Match
Best Local Similarity
Matches 20; Conserv
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Synthetic.
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                                                                        Sequence 20
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/note= "2'-O-methoxyethyl
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note= "2'-0-methoxyethyl (2'-MOE) gapmer"
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1.9e+02;
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                     The present invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding NAC, where the compound specifically hybridises with the nucleic acid molecule encoding NAC and inhibits the expression of NAC. The compound specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding NAC. Also described: (I) a composition comprising (I) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of NAC in cells or tissues comprising contacting the cells or tissues with (I); and (3) treating an animal having a disease or condition associated with NAC comprising administering (I) to the animal so that expression of NAC is inhibited (I) has cytostatic, nootropic, neuroprotective and antisinflammatory activities, and can be used in antisense therapy. The antisense compounds (I) are useful for modulating the expression of NAC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compounds, useful for modulating the expression of NAC for treating a disease or condition associated with the expression of
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      for treating a disease or
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/mod_base= OTHER
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/mod_base= OTHER
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   condition associated with
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Claim 3; Page 75; 147pp; English
                                                    New antisense compounds, useful for modulating the expression of NAC for treating a disease or condition associated with the expression of NAC, e.g. hyperproliferative disease or neurological disease.
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Synthetic.
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                                                                                                                                                                                                                                                                                    (ISIS-) ISIS PHARM INC.
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/mod_base= OTHER
/mote= "2'-O-methoxyethyl (2'-MOE)
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/mod_base= OTHER
/mote= "2'-0-methoxyethyl (2'-MOE)
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/mod_base= OTHER
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100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; cytostatic; nootropic; neuroprotective; antiinflammatory; antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming (EEM-11ke apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding NAC, where the compound specifically hybridises with the nucleic acid molecule encoding NAC and inhibits the expression of NAC. The compound specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding NAC. Also described: (1) a composition comprising (I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACC45206 standard; DNA; 20 BP
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                                                                                                                                                                                       modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human NAC
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                                                                    WO2003024988-A1
                                                                                                                                                                                                                                                                                      modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                             /*tag=
                                                                                                                                                                                                                                     /*tag= b
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                               /note= "phosphorothioate backbone"
                                                                                                                                                                                                                   note= "2'-0-methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                           mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       phosphorothicate oligonucleotide SEQ
                                                                                                                                                                                            . 20
                                                                                                                                              base= OTHER
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                                                                                                                  "2'-O-methoxyethyl (2'-MOE)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20
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27-MAR-2003

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ACC45154/c
ID ACC451
XX ACC451
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XX ACC451
XX Human:
XX Hu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound CC specifically hybridises with the nucleic acid molecule encoding NAC and CC inhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid CC molecule encoding NAC. Also described: (I) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC NAC in cells or tissues comprising contacting the cells or tissues with CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC antisfication; and can be used in antisense therapy. The CC antisense compounds (I) are useful for modulating the expression of NAC, e.g. hyperproliferative disease, neurological disease, or a disease CC and for treating a disease or condition associated with expression of CC useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or CC cman for acreeming protein (DEPCAP). NAC is located on human chromosome CC 17pl3. The present sequence represents a human NAC chimeric CC phosphorothioate antisense oligonucleotide, which is given in the CC exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local (
                                                                                                                            Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                       Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACC45154;
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                          modified_base
                                                                                                                                                                                                                                                     antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                          Human; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 6 A; 5 C; 5 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                       nootropic; neuroprotective; antiinflammatory
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Pred. No.
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1.9e+02;
hes 0; Indels
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RESULT 91
ACC45155/c
ID ACC451
XX
AC ACC451
XX
DT 16-JUN

ACC45155 standard;

DNA; 20

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ACC45155

16-JUN-2003

(first entry)

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20 GGGTTCTGAAAGCCCATTCC 1

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The present invention describes a compound (I) 8-50 nucleobases in length controlled to a nucleic acid molecule encoding NAC, where the compound specifically hybridises with the nucleic acid molecule encoding NAC and chihibits the expression of NAC. The compound specifically hybridises with the nucleic acid molecule encoding NAC and compound specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding NAC. Also described: (I) a composition comprising (I) complete acid molecule encoding NAC. Also described: (I) a composition comprising (I) composition and carrier or diluent; (2) inhibiting the expression of CC (I); and (3) treating an animal having a disease or condition associated with NAC comprising administering (I) to the animal so that expression of CC (I); and (3) treating administering (I) to the animal so that expression of CC (II); and compounds (I) are useful for modulating the expression of NAC, antiinflammatory activities, and can be used in antisense therapy. The compounds (I) are useful for modulating the expression of NAC, and for treating a disease, or condition associated with expression of CC (I); and for treating from aberrant apoptosis. The compounds are also CC (I); and season reason the proposis. The compounds are also cuseful as research reason to DEPCAD). NAC is also known as a death effector filament-forming CC (ED4-1); the apoptosis protein (DEPCAD). NAC is located on human chromosome Describioate antisense coligonucleotide, which is given in the corresponding to the present and prophylaxis.
                                                               Query Match
Best Local S
Matches 20
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                                                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                               exemplification of the present
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compounds, useful for modulating the expression of NAC for treating a disease or condition associated with the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-354583/33.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-SEP-2001; 2001US-00956712.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-SEP-2002; 2002WO-US029664.
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61 GGGTTCTGAAAGCCCATTCC
                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hyperproliferative disease or neurological disease.
                                                                                                                                                                                               BP; 5 A; 5 C; 6 G; 4 T; 0 U; 0 Other;
                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Preier
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/note= "2'.O-methoxyethyl (2'.MOE) gapmer"
16. .20
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/mod_base= OTHER
/note= "2'-O-methoxyethyl (2'-MOE) gapmer"
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                                                                                                 100.0%;
                                                                                                                                      0.4%;
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                                                                   <u>.</u>
                                                                                                    Score 20;
Pred. No.
   80
                                                                                                                                                                                                                                                                  invention
                                                                      Mismatches
                                                                                                                                   DB 1;
                                                                                                    1.9e+02
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                                                                                                                               Length 20
                                                                      Indels
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Human

NAC

chimeric phosphorothioate oligonucleotide SEQ

ID NO:15

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The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound CC specifically hybridises with the nucleic acid molecule encoding NAC and CC inhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid CC molecule encoding NAC. Also described: (I) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering the used in antisense therapy. The CC (I); and for treating administering the speciated with expression of CC (I); and for treating from aberrant apoptosis. The compounds are also CC (I); and for treating the expression of CC (I); and for treating the expression of CC (I); and for treating administering the expression of CC (I); and for treating from aberrant apoptosis. The compounds are also cuseful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or CC (I); and for treating the expression of CC (I); and for treating the expression of CC (I); and for treating the expression of CC (I); and the present sequence represents a human NAC (himeric (I); and the present sequence represents a human NAC (himeric (I); and the present sequence represents and the control of the presen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; cytostatic; nootropic; neuroprotective; antinflammatory; antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 15; Page 75; 147pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New antisense compounds, useful for modulating the expression of NAC for treating a disease or condition associated with the expression of NAC, e.g. hyperproliferative disease or neurological disease.
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Matches 20; Conserv
                       The present invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding NAC, where the compound specifically hybridises with the nucleic acid molecule encoding NAC and inhibits the expression of NAC. The compound specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding NAC. Also described: (1) a composition comprising (I) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of NAC in cells or tissues comprising contacting the cells or tissues with (I); and (3) treating an animal having a disease or condition associated with NAC comprising administering (I) to the animal so that expression of NAC is inhibited. (I) has cytostatic, nootropic, neuroprotective and antificial animal contactive and antifical animal contactive and animal contactive animal antifical animal contactive animal animal
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and for treating a disease or condition associated with expression of NAC, e.g. hyperproliferative disease, neurological disease, or a disease or diseorder arising from abservant apoptosis. The compounds are also useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. NAC is also known as a death effector filament-forming CED4-11ke apoptosis protein (DEPCAD). NAC is located on human chromosome 17p13. The present sequence represents a human NAC chimeric phosphorothicate antisense oligonucleotide, which is given in the exemplification of the present invention

antisense compounds (I) are useful for modulating the expression of NAC.

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New antisense compounds, useful for modulating the expression of NAC or for treating a disease or condition associated with the expression of NAC, e.g. hyperproliferative disease or neurological disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACC45198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          phosphorothicate antisense oligonuclectide, which is given exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CED4-like apoptosis protein (DEFCAP). NAC is located on human chromosome 17p13. The present sequence represents a human NAC chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 3; Page 75; 147pp; English
                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1121 AGGCTCCTGGGACCCAATGG 1140
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 AGGCTCCTGGGACCCAATGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 3 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                                                                  /*tag=
/mod_b
                                                                                                                                                                                                                                                       /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                 /mod_base= OTHER
/note= "2'-O-methoxyethyl (2'-MOB) gapmer"
                                                                                                                                                                                                                                                                                                                                                  note=
                                                                                                                                                                                                                                                                                                                                                                                   mod_base= OTHER
                                                                                                                                                                                                                       notes
                                                                                                                                                                                             6. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           phosphorothicate oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                          base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            describes a compound (I) 8-50 nucleobases in length
                                                                                                                                                                                                                                                                                                                                                  "phosphorothioate backbone'
                                                                                                                                                                                                                          -O-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 20; pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; L
1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                disease; apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ħ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NO:58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;

Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:26.

Human; cytostatic; nootropic; neuroprotective; antiinflammatory;

16-JUN-2003 ACC45166;

(first entry)

RESULT 93 ACC45166/c

ACC45166 standard;

DNA;

20 ΒP 밁 S

20

AGCTGAAGGAGTTCCAGCTT 1

581 AGCTGAAGGAGTTCCAGCTT 600

Query Match Best Local S Matches 20

Similarity

20;

Conservative

0.4%; 5--100.0%; Pred. No. . 179 0; Mismatches

DB 1; Lo

Length 20 Indels

0

Gaps

0

Sequence 20 BP; 5 A; 6 C; 4 G; 5 T; 0 U; 0 Other;

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RESULT 95
ACC45202(c
ID ACC45
XX ACC45
XX ACC45
XX ACC45
XX Humar

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound CC specifically hybridises with the nucleic acid molecule encoding NAC and CC inhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid CC molecule encoding NAC. Also described: (1) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comporising administering (I) to the animal so that expression of CC (I); and (3) treating an animal having a disease or condition associated CC with NAC compounds (I) has cytostatic, nootropic, neuroprotective and CC antisense compounds (I) are useful for modulating the expression of CC (I); and (I) the cytostatic, nootropic, neuroprotective and CC antisense compounds (I) are useful for modulating the expression of CC (I); and (I) are useful for modulating the expression of CC (I); and (I); and (I) are useful for modulating the expression of CC (I); and (I); and (I); are useful for modulating the expression of CC (I); and (I); and (I); are useful for modulating the expression of CC (I); and (I); are useful for modulating the expression of CC (I); and (I); are useful for modulating the expression of CC (I); and (I); are useful for modulating the expression of CC (I); and (I); are animal animal (I); are animal animal (I); are a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                             antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACC45202 standard;
                                                                            Synthetic.
                                                                                                                    Homo sapiens
                                                                                                                                                                                                 phosphorothicate;
                                                                                                                                                                                                                                                                                                                                                     Human; cytostatic; nootropic; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human NAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  phosphorothicate antisense oligonuclectide, which exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compounds, useful for modulating the expression of NAC for treating a disease or condition associated with the expression of NAC, e.g. hyperproliferative disease or neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-354583/33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-MAR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-SEP-2001; 2001US-00956712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4565 GAGTGACATCGGTGGAGGTT 4584
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                              chimeric phosphorothioate oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 5 A; 9 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Freier SM;
    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>.</u>
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Pred. No. 1.9e+0
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                                                                                                                                                                                                                                                                                                                                                                                                                                  NO:62
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RESULT 96 ACC45159/c

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ACC45159;

ACC45159 standard; DNA; 20 BP

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                                                                                                                                                   CC targeted to a nucleic acid molecule encoding NAC, where the compound compectable to a nucleic acid molecule encoding NAC and cinhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding NAC. Also described: (1) acomposition comprising (1) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC NAC in cells or tissues comprising contacting the cells or tissues with CC (1); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (1) to the animal so that expression of CC NAC is inhibited. (1) has cytostatic, nootropic, neuroprotective and CC antisense compounds (1) are useful for modulating the expression of NAC, and for treating a disease or condition associated CC and for treating a disease or condition associated with expression of CC useful as research reagents and kits, or for diagnostics, therapeutics CC and prophylaxis, e.g. to prevent or delay infection, inflammation or customar formation. NAC is also known as a death effector filament-forming CC CED4-like apoptosis protein (DEFCAP). NAC is located on human chromosome CC phosphorothioate antisense oligonucleotide, which is given in the CC exemplification of the present invention
                                                            Query Match
Best Local S
Matches 20
                                                                                                                          Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a compound (I) 8-50 nucleobases in length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15; Page 76; 147pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-354583/33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO2003024988-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antisense compounds, useful for modulating the expression of NAC treating a disease or condition associated with the expression or e.g. hyperproliferative disease or neurological disease.
                              4789 GGACTCCTGCCACTCAGCAG 4808
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                                                          20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CF,
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                                                                             Similarity
                                                             Conservative
                                                                                                                          BP; 3 A; 5 C; 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note=
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                                                       100.0%; ***
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          base= OTHER
                                                                         0.4%; Score 20;
100.0%; Pred. No
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "2'-O-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a
                                                                                                                           G; 4 T; 0 U; 0 Other;
                                                             red. No. 1.
                                                          1.9e+02;
0;
                                                                                            DB 1; Length 20
                                                               0
                                                               Gaps
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Claim 3; Page 75; 147pp;
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                                                                                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; cytostatic; nootropic; neuroprotective; antinflammatory; antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                    WO2003024988-A1
                                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                            (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-JUN-2003
                                                                                                                                                                                                                                2003-354583/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                             sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAC
                                                                                                                                                                                                    hyperproliferative disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimeric phosphorothioate oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                              Freier SM;
                                                                                                                                                                                                                                                                                                                                               16. .20
/*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                              /note=
                                                                                                                                                                                                                                                                                                                                                                      mod.
                                                                                                                                                                                                                                                                                                                                                                                         /note=
                                                                                                                                                                                                                                                                                                                                                                            *tag=
                                                                                                                                                                                                                                                                                                                                                                                                  mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                              base= OTHER
e= "2'-0-methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                  _base= OTHER
== "2'-O-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                                                                                                                                                                                                                                 ი
                                                                                                                                                                                                                                                                                                                                                                                          "phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                          ρ
                                                                                                                                                                                      English.
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R
                                                                                                                                                                                                   neurological disease
                                                                                                                                                                                                                                                                                                                                                                                           backbone*
                                                                                                                                                                                                                                                                                                                                                              gapmer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ID NO:19
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The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound competifically hybridises with the nucleic acid molecule encoding NAC and controlled in the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid comolecule encoding NAC. Also described: (1) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of NAC in cells or tissues comprising contacting the cells or tissues with CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC NAC is inhibited. (I) has cytostatic, nootropic, neuroprotective and CC antisense compounds (I) are useful for modulating the expression of CC NAC, and for treating a disease or condition associated with expression of CC NAC, e.g. hyperproliferative disease, neurological disease, or a disease CC or disorder arising from abservant apoptosis. The compounds are also cuseful as research respents and kits, or for dispnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or CC tumour formation. NAC is also known as a death effector filament-forming CC 17p13. The present sequence represents a human NAC chimeric CC phosphorochioate antisense oligonucleotide, which is given in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antisense compounds, useful for modulating the expression of NAC treating a disease or condition associated with the expression of
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    RESULT 97
ACC45167/c
ID ACC4511
XX ACC451
XX ACC451
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XX Human
DH Human
XX Homo 8
OS Synthe
YT modifi
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XX EY
XX WEV
XX WEV
XX 19-SE
XX WEV
XX ISIS

The present invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding NAC, where the compound specifically hybridises with the nucleic acid molecule encoding NAC and inhibits the expression of NAC. The compound specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding NAC. Also described: (I) a composition comprising (I) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of NAC in cells or tissues comprising contacting the cells or tissues with (I); and (3) treating an animal having a disease or condition associated with NAC comprising administering (I) to the animal so that expression of
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Sequence 20 BP; 6 A; 4 C; 8

G; 2 T; 0 U; 0 Other;

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Query Match
Best Local S
Matches 20
                              New antisense compounds, useful for modulating the expression of NAC for treating a disease or condition associated with the expression of NAC, e.g. hyperproliferative disease or neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; cytostatic; nootropic; neuroprotective; antiinflammatory; antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human NAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACC45167 standard; DNA; 20
Claim 3; Page 75; 147pp; English.
                                                                                                                                                                         (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                          19-SEP-2001; 2001US-00956712.
                                                                                                                                                                                                                                            19-SEP-2002; 2002WO-US029664.
                                                                                                                                                                                                                                                                                                               WO2003024988-A1
                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neurological disease; infection; inflammation; tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACC45167
                                                                                                      2003-354583/33.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                       Freier
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/mod_base= OTHER
/note= "2'-O-methoxyethyl (2'-MOB) gapmer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                   /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          pom/
                                                                                                                                                                                                                                                                                                                                                   note=
                                                                                                                                                                                                                                                                                                                                                                     mod.
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100.0%; Pr.
                                                                                                                                                                                                                                                                                                                                                 base= OTHER
== "2'-O-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          _base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 20; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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Best Loc
Matches
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 New antisense compounds, useful for modulating the expression of NAC
                                  WPI; 2003-354583/33
                                                                                                                                                                                                                            WO2003024988-A1.
                                                                                                                                                                                                                                                                                                      modified_base
                                                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ACC45175;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACC45175 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 5 A; 4 C; 2 G; 9 T; 0 U; 0 Other;
                                                                Bennett CF, Freier SM,
                                                                                                                             19-SEP-2001; 2001US-00956712
                                                                                                                                                             19-SEP-2002; 2002WO-US029664.
                                                                                                                                                                                              27-MAR-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; cytostatic; nootropic; neuroprotective; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local
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                                                                                                ISIS
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                                                                                                PHARM INC.
                                                                                                                                                                                                                                                                                           /*tag= b
/mod_base= OTHER
/note= ^2'-O-methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                         /note=
                                                                                                                                                                                                                                                                                                                                                                                                     /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                      'mod_base= OTHER
'note= "2'-0-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.4%; Sur
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                     _base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                      "phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
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Pred. No. 1.9e+
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                      backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
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                                                                                                                                                                                                                                                                                                                       gapmer"
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밁 S

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for treating a disease or condition associated with the expression of NAC, e.g. hyperproliferative disease or neurological disease.
147pp; English
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The present invention describes a compound (I) 8-50 nucleobases in length crargeted to a nucleic acid molecule encoding NAC, where the compound CC specifically hybridises with the nucleic acid molecule encoding NAC and cinhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid CC molecule encoding NAC. Also described: (1) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC NAC in cells or tissues comprising contacting the cells or tissues with CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC NAC is inhibited. (I) has cytostatic, nootropic, neuroprotective and CC antisense compounds (I) are useful for modulating the expression of NAC, e.g. hyperproliferative disease, neurological disease, or a disease or condition associated with expression of CC useful as research reagents and kits, or for diagnostics, therapeutics CC and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. NAC is also known as a death effector filammation or CC CED4-like apoptosis protein (DEFCAP). NAC is located on human chromosome CC phosphorothioate antisense oligonucleotide, which is given in the campulation of the present sequence represents a human NAC chimeric CC phosphorothioate antisense oligonucleotide, which is given in the

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Matches
                                                         Query Match
Best Local (
                                                                                            Sequence 20
                       2929 GTCCTCAAGGTCACCAGAAA 2948
20
                                           20;
                                                         Similarity
                                             ilarity 100.0%; Pred. No. 1.9
Conservative 0; Mismatches
                                                                                            BP; 3 A; 4 C; 6 G; 7 T; 0 U; 0 Other;
                                                                       0.4%;
                                                        Score 20;
Pred. No.
                                                                     DB 1; Length 20
                                                          1.9e+02;
                                               0,
                                              Gaps
                                                0,
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밁 S

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RESULT 99
ACC45188/c
                                                                                           Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:48.
                                                                                                   16-JUN-2003
                                                                                                                  ACC45188 standard; DNA; 20 BP
                                                                                                  (first entry)
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Human; cytostatic; nootropic; neuroprotective; antiinflammatory; antisense therapy, NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation; modified_base Synthetic. modified_base modified_base Homo sapiens. phosphorothicate; /note= " Location/Qualifiers note= /*tag= a
/mod_base= OTHER _base= OTHER e= "2'-O-methoxyethyl (2'-MOE) "phosphorothioate backbone" σ

note= "2'-O-methoxyethyl (2'-MOE) gapmer

base= OTHER

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RESULT 100
ACC45165/c
ID ACC451
XX
ACC451
XX
AC ACC451
XX
DT 16-JUN
XX
DE Human
XX
Human
XX
KW Human;
KW antise
KW death
KW neurol
KW phosph
XX
OS Homo s
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound compound the present of the nucleic acid molecule encoding NAC and compound the expression of NAC. The compound specifically hybridises with the nucleic acid molecule encoding NAC and compound specifically hybridises with the teast an 8-nucleobase portion of an active site on a nucleic acid molecule encoding NAC. Also described: (1) a composition comprising (I) composition comprising (I) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of (C) (I); and (3) treating an animal having a disease or condition associated (I); and (3) treating an animal having a disease or condition associated comprising administering (I) to the animal so that expression of (I) and (3) treating an animal having a disease or condition associated (I) has cytostatic, notropic, neuroprotective and (I) and the compounds (I) are useful for modulating the expression of (I) and for treating a citivities, and can be used in antisense therapy. The candition associated with expression of (I) and for treating a disease or condition associated with expression of (I) and for treating a disease or condition associated with expression of (I) and for treating from abservant apoptosis. The compounds are also (I) as a second condition associated with expression of (I) as a constant apoptosis. The compounds are also (I) as a constant apoptosis. The compounds are also (I) as a constant apoptosis. The compounds are also (I) and prophylaxis, e.g. to prevent or delay infection, inflammation or constituent formation or the present sequence represents a human NAC chimeric (I) phis present sequence oligonucleotide, which is given in the compounds antisense oligonucleotide, which is given in the
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Best Local 9
                                                                                   Human; cytostatic; nootropic; neuroprotective; antiinflammatory; antiense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector fillament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                       ACC45165
                                                                                                                                                                                                                                                                                                                               ACC45165 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 3; Page 75; 147pp; English
  Homo sapiens
Synthetic.
                                                                                                                                                                                              Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New antisense compounds, useful for modulating the expression of NAC for treating a disease or condition associated with the expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bennett CF,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Freier SM;
                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; 3 C; 8
                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.4%; ET
100.0%; Pr
'--- 0;
                                                                                                                                                                                                                                                                                                                               20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 20; DB 1;
Pred. No. 1.9e+(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
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                                                                                                                                          CC specifically hybridises with the nucleic acid molecule encoding NAC and cinhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid collecule encoding NAC. Also described: (1) a composition comprising (1) composition comprising (1) composition comprising (1) comparison of the cells or tissues comprising contacting the cells or tissues comprising contacting the cells or tissues with CC (1); and (3) treating an animal having a disease or condition associated comprising administering (1) to the animal so that expression of NAC is inhibited. (1) has cytostatic, nootropic, neuroprotective and CC nactisficating activities, and can be used in antisense therapy. The CC and for treating a disease or condition associated with expression of NAC, e.g. hyperproliferative disease, neurological disease, or a disease or disorder arising from aberrant apoptosis. The compounds are also cuseful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or CC tumour formation. NAC is also known as a death effector filament-forming CCD4-like apoptosis protein (DEFCAP). NAC is located on human chromosome components approach antisense oligonucleotide, which is given in the components of the present sequence represents a human NAC chimeric components and the present invention.
                                                                                     Query Match
Best Local 9
                                                                     Matches
                                                                                                                                          Sequence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a compound (I) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding NAC, where the compound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 3; Page 75; 147pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense compounds, useful for modulating the expression of NAC or for treating a disease or condition associated with the expression of
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RESULT 101 ACC45177/c ID ACC451

ACC45177 standard; DNA; 20 BP

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The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound CC specifically hybridises with the nucleic acid molecule encoding NAC and CC inhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid CC molecule encoding NAC. Also described: (1) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC NAC in cells or tissues comprising contacting the cells or tissues with CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC NAC is inhibited. (I) has cytostatic, nostropic, neuroprotective and CC antisnflammatory activities, and can be used in antisense therapy. The CC and for treating a disease or condition associated with expression of NAC, C and for treating a disease or condition associated with expression of CC NAC, e.g. hyperproliferative disease, neurological disease, or a disease CC or disorder arising from aberrant apoptosis. The compounds are also CC useful as research reagents and kits, or for diagnostics, therapeutics CC and prophylaxis, e.g. to prevent or delay infection, inflammation or CC cumour formation. NAC is also known as a death effector filament-forming CC CED4-like apoptosis protein (DEFCAP). NAC is located on human chromosome cromation account of the protein (DEFCAP) and the protein consonne co
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17p13. The present sequence represents a human NAC chimeric phosphorothioate antisense oligonucleotide, which is given exemplification of the present invention
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specifically hybridises with the nucleic acid molecule encoding NAC and inhibits the expression of NAC. The compound specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding NAC. Also described: (1) a composition comprising (1) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of NAC in cells or tissues comprising contacting the cells or tissues with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              note= "phosphorothicate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nootropic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  _base= OTHER
e= "2'-O-met
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "2'-O-methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -O-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 20;
; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 6 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antiinflammatory;
                                                                                                                             where the compound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gapmer'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease; apoptosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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RESULT 103
ACC45199/c
ID ACC451
XX ACC451
XX ACC451
XX Inman
DE Human
XX Human;
KW Antise
KW Antise
KW Ineurol
Ineur
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and for treating a disease or condition associated with expression of NAC, e.g. hyperproliferative disease, neurological disease, or a disease or disorder arising from aberrant apoptosis. The compounds are also useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflammation or tumour formation. NAC is also known as a death effector filament-forming CED4-like apoptosis protein (DEPCAP). NAC is located on human chromosome 17p13. The present sequence represents a human NAC chimeric phosphorothicate antisense oligonucleotide, which is given in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; cytostatic; nootropic; neuroprotective; antiinflammatory; antisense therap; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-JUN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACC45199;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACC45199 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (I); and (3) treating an animal having a disease or condition associated with NAC comprising administering (I) to the animal so that expression of NAC is inhibited. (I) has cytostatic, nootropic, neuroprotective and antiinflammatory activities, and can be used in antisense therapy. The antisense compounds (I) are useful for modulating the expression of NAC,
   WPI; 2003-354583/33
                                                               Bennett CF, Freier SM;
                                                                                                                                                                                                                                                                                                                                                                            WO2003024988-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
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                                                                                                                                                                                      19-SEP-2001; 2001US-00956712
                                                                                                                                                                                                                                                  19-SEP-2002; 2002WO-US029664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3846 CCGCTGGCCCAACACGGGTC 3865
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                                                                                                                            SISI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                            PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mod_base= OTHER
/note= "2'-O-methoxyethyl (2'-MOB) gapmer"
16. .20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        *tag= b
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100.0%; Pr/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  _base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                  "2'-O-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ი
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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New antisense compounds, useful for modulating the expression of NAC or for treating a disease or condition associated with the expression of NAC, e.g. hyperproliferative disease or neurological disease.
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Example 15; Page 76; 147pp; English.

The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound CC specifically hybridises with the nucleic acid molecule encoding NAC and C inhibits the expression of NAC. The compound specifically hybridises with CC at least an 8-nucleobase portion of an active site on a nucleic acid CC molecule encoding NAC. Also described: (I) a composition comprising (I) CC and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC (I); and (3) treating an animal having a disease or condition associated CC with NAC comprising administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I) to the animal so that expression of CC (I); and for treating administering (I); and the expression of CC (I); and the compounds (I) are useful for modulating the expression of CC (I); and for treating administering (I); and the expression of CC (I); and for treating administering (I); and the expression of CC (I); and the Sequence 20 BP; 6 A; 9 C; 2 G; 3 T; 0 U; 0 Other;

S Query Match
Tocal Similarity Matches 20; 4575 GGTGGAGGTTGTCTTGGACA 4594 Conservative 0.4%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 1.9e+02; 0, Mismatches Indels 0 Gaps 0

문

20

RESULT 104
ADP11640/c
ID ADP116
XX ADP116
XX IT-AUG
XX I Taqman probe of the invention #323. 12-AUG-2004 ADP11640 ADP11640 standard; DNA; 20 BP (first entry)

transplant rejection; immune system; rheumatoid arthritis; inflammatory bowel disease; multiple sclerosis; HIV; AIDS; lupus; ss; probe.

Homo sapiens.

WO2004042346-A2

24-APR-2003; 2003WO-US012946.

24-APR-2002; 2002US-00131831. 20-DEC-2002; 2002US-00325899.

(EXPR-) EXPRESSION DIAGNOSTICS INC.

Wohlgemuth J, Rosenberg S; Fry K, Woodward R, Ly N, Prentice Ģ Morrie M;

WPI; 2004-400724/37

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RESULT 105
ADP10980
ID ADP109
XX ADP109
XX ADP109
XX Set 1
XX Uransp
XX Inflam
XX UW02004
XX W02004
XX UC2004
XX UC2006
XX UC2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, xenotransplant rejection or mechanical organ replacement rejection, in a individual. The method is also useful in assessing the immune status of an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lungs, inflammatory bowel diseases, multiple sclerosis, HTV/AIDS or viral, bacterial or fungal infection. The present sequence represents a Tagman probe for a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft rejection and other disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rejection, e.g. cardiac or kidney transplant rejection, in an comprises detecting the expression level of one or more genes. methods, system and kits are useful in diagnosing or monitoring
                                                                                                                                                                                                                      Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transplant rejection; immune system; rheumatoid arthritis; inflammatory bowel disease; multiple sclerosis; HIV; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set 1 left PCR primer for marker probe #325.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-400724/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wohlgemuth J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-APR-2002;
20-DEC-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 58; SEQ ID NO 1649; 1762pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (EXPR-) EXPRESSION DIAGNOSTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4970 TGCCAGGGATGCCACAGGGG 4989
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2002US-00325899.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Fry K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Woodward R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Prentice
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ss; primer.
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                                              comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in a xenotransplant rejection or mechanical organ replacement rejection, in a individual. The method is also useful in assessing the immune status of an individual The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lupus, inflammatory bowel diseases, multiple sclerosis, HIV/AIDS or viral, bacterial or fungal infection. The present sequence represents a primer for a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft rejection and other disorders.
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Sequence 20
  BP; 2 A; 4 C; 6 G; 8 T; 0 U; 0 Other;
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Query Match Best Local S Matches 20 20; Similarity 0.4%; ilarity 100.0%; Conservative (0, Score 20; Pred. No. Mismatches DB 1; L Length Indels 20 0, Сарв <u>,</u>

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                  4945
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              GCCTTTGCTGGGCATTATGT 4964
 GCCTTTGCTGGGCATTATGT 20
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2136/c
ADP12136
standard;
DNA;
 20
 ВP
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ADP12136;

Set 2 right 12-AUG-2004 PCR primer for marker probe #242 (first entry)

eapiens.

WO2004042346-A2

inflammatory transplant

rejection; immunery bowel disease;

immune system; rheumatoid
sease; multiple sclerosis;

arthritis; HIV; AIDS;

88; lupus;

primer

21-MAY-2004

24-APR-2003; 2003WO-US012946

24-APR-2002; 2002US-00131831. 20-DEC-2002; 2002US-00325899.

(EXPR-) EXPRESSION DIAGNOSTICS INC

Wohlgemuth J, Rosenberg S; Fry K, Woodward R, Ł z Prentice J, Morris

WPI; 2004-400724/37.

Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of

Claim 58; SEQ ID NO 2145; 1762pp; English.

RESULT 106
ADP12136/6
ADP12136/6
ADP12136/6
ADP12136/6
AC ADP123
AC ADP121
AC AC AC AC
AC AC AC
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AC AC The present invention relates to diagnosing or monitoring transplant rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in ar xenotransplant rejection or mechanical organ replacement rejection, in ar individual. The method is also useful in assessing the immune status of an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lupus, inflammatory bowel diseases, multiple sclerosis, HIV/AIDS or 21

The present rejection, 6

t invention relates to diagnosing or monitoring transplant e.g. cardiac or kidney transplant rejection, in an indivi-

individual

Claim

58; SEQ ID NO 989; 1762pp; English

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RESULT 107
ADP1131/c
ID ADP113
XX ADP113
XX ADP113
XX Transp
XX Uransp
XX Uransp
XX Uransp
XX Uransp
XX Uransp
XX Uransp
XX HOMO 8
XX WO2004
XX 21-MAY
XX 24-APR
PP 24-APR
PR 20-DEC
XX Wohlge
PI Rosenb
XX Well; 2
XX Well; 2
XX Uransp
XX Uransp
XX Transp
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     Query Match
Best Local Similarity
Matches 20; Conserv
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                       The present invention relates to diagnosing or monitoring transplant rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in ar individual. The method is also useful in assessing the immune status of an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthitis, lung, inflammatory bowel diseases, multiple sclerosis, HIV/AIDS or viral, bacterial or fungal infection. The present sequence represents a primer for a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft rejection and other disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Diagnosing or monitoring transplant rejection, e.g. heart, kidney, lipancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 58; SEQ ID NO 1320; 1762pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-400724/37.
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20-DEC-2002; 2002US-00325899.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      transplant rejection; immune system; rheumatoid arthritis; lupus; inflammatory bowel disease; multiple sclerosis; HIV; AIDS; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Set 1 right PCR primer for marker probe #325.
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           Conservative
                                                                                                                       BP;
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100.0%; **
                                                                                                                       6 C;
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100.0%; Pred. No.
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                                                                                                                       5 G; 4 T; 0 U; 0 Other;
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                                Score 20;
Pred. No.
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           Mismatches
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1.9e+02;
     DB 1; Length 20;
1.9e+02;
hes 0; Indels
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RESULT 108
ADP11888
ID ADP118
XX ADP118
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DT 12-AUG
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XX HOMO 8
XX HOMO 9
XX HOMO 121-MAY
XX 21-MAY
XX 24-APF
PR 20-DE(
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                                                                                                                    Matches
                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 8 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-400724/37
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20-DEC-2002; 2002US-00325899.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transplant rejection; immune system; rheumatoid arthritis; inflammatory bowel disease; multiple sclerosis; HIV; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set 2 left PCR primer for marker probe #240.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 58; SEQ ID NO 1897; 1762pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO2004042346-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (EXPR-) EXPRESSION DIAGNOSTICS INC.
                                                                                                                                              Local
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                                                            4825 AGCCCTTGACCCTTGAGTCC 4844
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                                                                                                                 20;
                                                                                                                                                    Similarity
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                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7
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                                                                                                                    Mismatches
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                                                                                                                                                                             DB 1;
                                                                                                                                                 1.9e+02;
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                                                                                                                 Gaps
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RESULT 109 ACK28659 ID ACK286

ACK28659 standard; DNA; 25 BP.

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RESULT 110
AAH91641/c
ID AAH916:
XX
AC AAH916:
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                                                                                                                                                                                                      CC perfect match, perfect mismatch, antisense mitherse mismatch.

CC Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, in analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis computed for analysis comprises comparises and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring gene expression levels, identifying biallelic markers or polymorphisms, or family members of a gene and a cross-species comparison. Each of the nucleic acid strother comprises a tag sequence. The array of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dotable hybridisation to identify or detect the sequence or specific mutations of any gene, in mapping the 5' termini of mRNA molecules by comparisons with the sequence or in screening cDNA or genomic libraries or subclones containing segments of DNA that have been concleic acid probes incorporated in the microarray. Note: The sequence contain for this patent can also be obtained in electronic format directly from USPTO at segdata.uspto.goc/sequence.html
                                                                                                                                     Query Match
Best Local S
Matches 21
  AAH91641
                          AAH91641 standard; DNA; 28 BP.
                                                                                                                                                                                      Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 128640; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence or specific mutations of any gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New array of nucleic acid probes, useful for Southern, Northern or dot-blot hybridization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-567953/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-MAR-2001; 2001US-0276759P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-MAR-2002; 2002US-00098263
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cross-species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EST; ss; probe; expressed sequence tag; microarray; gene expression;
genetic variation; biallelic marker; polymorphism; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human microarray DNA oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (AFFY-) AFFYMETRIX INC
                                                                                                                                                Local Similarity
                                                                                                            3444 GGAGCAGGAGAAACCTCAGCTGC 3466
                                                                                                                                     21;
                                                                                       w
                                                                                    GGAGCAGGAGAATCCTCAGCTAC 25
                                                                                                                                      Conservative
                                                                                                                                                                                    BP; 8 A; 6 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  comparison.
                                                                                                                                                0.4%;
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                                                                                                                                                Score 19.8; DB 1
Pred. No. 2.1e+02
                                                                                                                                      Mismatches
                                                                                                                                                              DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                situ hybridization, in identify or detect the
                                                                                                                                                            Length 25;
                                                                                                                                      Indels
                                                                                                                                     0,
                                                                                                                                     Gaps
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 RESULT 111
AAA57855
ID AAA578
XX
XX
AC AAA578
XX
DT 11-OCT
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DE DEOXY-
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W Ribozy
KW Ribozy
KW Ruclei
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1;
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                                                                                                                                                                                                                                                                         Local
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The present invention describes a method for detecting the presence of polymorphisms associated with inflammatory bowel diseases such as ulcerative colitis and Crohn's disease. The methods can be used to detect the presence of genetic polymorphisms associated with inflammatory bowel disease and correlating their occurrence with disease states. They may be used in this way for phenotypic correlations, forensics, paternity testing, medicine and genetic analysis. The present sequence is a polymorphic site described in the exemplification of the invention
Ribozyme; catalytic RNA; analyte detection; nucleic acid substrate; in vitro selection; conformation dependent activity; allosteric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; inflammatory bowel disease; Crohn's disease; ulcerative colitis; single nucleotide polymorphism; SNP; chromosome 19p13; paternity test; chromosome 5q31-33; forensic test; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 28 BP; 3 A; 9 C; 2 G; 13 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Testing for the presence of polymorphisms associated with inflammatory bowel disease, using a hybridization assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human inflammatory bowel disease associated polymorphic site #716
                                                                                                Deoxy-A22-tagged substrate oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-367874/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10-DEC-1999; 99US-0170257P.
10-APR-2000; 2000US-0196046P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 11-DEC-2000; 2000WO-US033632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200142511-A2
                                                                                                                                                                                                                                                             AAA57855 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                            1177 ATCAGAGAAAGAGAGAGAGAAA 1200
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BLLIPSIS BIOTHERAPEUTICS CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 69; 463pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hudson TJ,
                                                                                                                                                                                                                                                                                                                                                                                              ATCAGAGAGANAGAGAGAGAGA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.4%;
nilarity 87.5%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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/note= "SNP, optionally insertion or deletion at this
                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lander ES,
                                                                                                                                                                                                                                                               28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 19.8;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2.2e+02;
3;
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                        effector molecule;
ribozyme ligase;
     activation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 28;
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CC for the catalytic activity of the nucleic acid sequence; and contacting conditions allowing catalytic activity of nucleic acid sequences. The CC catalytic nucleic acid sequence will be able to convert the nucleic acid sequences. The CC catalytic nucleic acid sequence will be able to convert the nucleic acid sequences. The CC present. The nucleic acid product only if the analyte of interest is CC present. The nucleic acid catalytic product is then amplified, and a CC significant increase in the amount of product indicates the presence of the analyte in the sample. The method is useful for the qualitative or CC assays. The invention describes the in vitro selection of a ribozyme CC ligase (L1; AAA57859, AAA57860) which is catalytically active only in the presence of an oligonuclectide effector (AAA57854). The L1 ribozyme CC indicate was selected from a pool of RNA molecules comprising a central CC constant sequence regions (the N90 RNA pool; AAA57851). In the presence CC these ffector, selection was performed using one of the tagged CC substrate molecules AAA57855-A57857. RNAs with ligase activity (i.e., those which have become ligated to the substrate molecule) were reverse CC transcribed using the effector oligo, and then PCR amplified using the effector oligo, and then PCR amplified using the catalyte). In the absence of the substrate used for the selection A ribozyme ligase, L1, was selected via this procedure. L1 cancive conformation (AAA57859) in the presence of the effector oligo (analyte). In the absence of the effector, L1 adopts an catalyte conformation (AAA57859) in the presents the CC deoxy-A22-tagged substrate oligonucleotide. The dA22 tag enables catalytes and the present sequence is not given in the specification, but is created from the inforamtion given on page 11
                                                                     Query Match
Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting an analyte in a sample comprises providing nucleic acid sequence which is catalytically active in presence of analyte, contacticatalytic nucleic acid with substrate and amplifying catalytic product.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a method of detecting an analyte in a sample. The method comprises providing a nucleic acid sequence which is initially catalytically inactive, but which becomes catalytically active in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page; 36pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nathan A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 misc_binding
                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               presence of an analyte (the effector); providing a nucleic acid substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (INTE-) INTELLIGENE LTD
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5394 AAAAAATACAAAAAAGAAAAAATGAA 5419
                                                                                                                Similarity
                                                                                                                                                                                                                    28 BP;
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/*tag= a

24. .28

/*tag= b

/*tound_moiety= "Bases 13-17 of N90 RNA pool (AAA57851)"
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                                                                                                                                                                                                                    A; 2 C; 1 G; 0 T; 2 U; 0 Other;
                                                                                                        0.4%;
                                                                         1; Mismatches
                                                                                                            Score 19.6; DB 1;
Pred. No. 2.3e+02;
                                                                                                                                                  Length 28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to regulatable, catalytically active nucleic acids (RCANAs) which are regulated by polypeptides. These a useful for regulating gene expression, in assays for detecting the presence of ligands, for activation of an effector of RCANA, and therapy. The present sequence is an oligonucleotide substrate used construction of an RCANA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ellington AD, hear-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New regulatable, catalytically active nucleic acids (RCANA), useful in gene therapy (particularly for regulating gene expression), or in assays for detecting the presence of ligands or activation of an effector of
  regulatable catalytically active nucleic acid; RCANA; catalytic domain; regulation; screening; gene therapy; biological pathway regulation; regulatory element; metabolic pathway; ribozyme; ss.
                                                                                                                   Substrate RNA related oligonucleotide SEQ ID NO:25
                                                                                                                                                                         20-NOV-2003
                                                                                                                                                                                                                                ADA39569;
                                                                                                                                                                                                                                                                                    ADA39569 standard; RNA; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 28 BP; 23 A; 2 C; 1 G; 0 T; 2 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Regulatable, catalytically active nucleic acid substrate #1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 6; Page 75; 126pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RCANA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200196559-A2
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                                                                                                                                                                         (first entry)
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/note= "5' biotinylated"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 19.6; D
Pred. No. 2.3e
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 28;
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RESULT 114
ADQ96960
ID ADQ969
XX ADQ969
XX ADQ969
XX ADQ969
DT 23-SEP
XX RCANA,
KW RCANA,
KW indust
XX Uniden
XX Uniden
XX Uniden
XX US2004
PD 01-UUL
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Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           its coding potential and a regulatory domain that recognises an effector that alters the function of the catalytic domain, and contacting the regulatory domain with an effector to regulate production of the product. Also described: (1) regulating a biological pathway in cell; and (2) screening a population of cells for a cell that produces a bioproduct. The methods are useful for regulating a biological pathway in cell, or regulating production of a product in a cell. The RCANAs are useful as regulatory elements to control the expression of genes in a metabolic pathway, or as regulated selectable markers to increase a selective pressure favouring or disfavouring production of a targeted bioproduct. The RCANAs are also useful for in vitro or in vivo sensing or detection, and in gene therapy. The present sequence represents an RNA substrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   product in a cell. The method comprises inserting a regulatable catalytically active nucleic acid (Rohn into a gene that produces the product or regulates the production of the product in the cell, where the RCANA comprises a catalytic domain which modifies a transcript to alter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Regulating production of a product in a cell, comprises inserting a regulatable catalytically active nucleic acid into a gene that produces the product or regulates the production of the product in the cell.
                                                                                                                            RCANA; catalytically active regulatable nucleic acid; ss; ribozyme; aptamer; effector domain; nucleic acid catalyst domain; gene therap; industrial biosynthesis; bioremediation; Ribosomal Ll ligase.
                                                                                                                                                                                                                                           23-SEP-2004
                                                                                                                                                                                                                                                                                                                  ADQ96960 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 6; Page 76; 128pp; English.
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                                                         US2004126882-A1
                                                                                            Unidentified
                                                                                                                                                                                                     Ribozyme substrate oligonucleotide S28A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oligonucleotide, which is used in an example from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              present invention describes a method for regulating production of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                  DNA;
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Pred. No. 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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15-JUN-2000; 2000US-0212097P.
14-SEP-2000; 2000US-006661658.
20-SEP-2000; 2000US-00666870.
14-JUN-2001; 2001US-00883119.
24-SEP-2001; 2001US-0324715P.
                                                                                                                                                                                                                                                                                            24-SEP-2002; 2002US-00254568.
                                                                                                                                                                                                        Davidson E,
                                                                                                                                                                                                            Ellington AD,
                                                                                                                                                                                                                                    (DAVI/)
                                                                                                                                                                                                                                 RIED/
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                                                                                                                                                                                                                    COX J C.
RIEDEL T.
WILSON C.
CLOAD S T.
KEEFE A D.
                                                                                                                                                                                                                                       DAVIDSON E.
                                                                                                                                                                                                                                                       ELLINGTON A
HESSELBERTH
                                                                                                                                                                                                                                                ROBERTSON M P.
                                                                                                                                                                                                                                                     THOMPSON K.
                                                                                                                                                                                                        Cox JC,
                                                                                                                                                                                                            Hesselberth
                                                                                                                                                                                                                                                        J.
                                                                                                                                                                                                        herth J, T
Riedel T,
                                                                                                                                                                                                         Thompson Wilson
                                                                                                                                                                                                         O X
                                                                                                                                                                                                         Robertson MP,
Cload ST, Ke
                                                                                                                                                                                                         Keefe AD;
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Novel regulatable, catalytically active nucleic acid comprising effector domain, and catalyst domain which comprises randomized catalytic residues and is regulated by effector that interacts with effector domain.

Example 6; SEQ ID NO 25; 78pp; English.

cd domain in which one or more critical catalytic residues of the nucleic catalytic domain are regulated by an effector that interacts with the catalytic domain. Also included are a nucleic acid comprising a gene, a ccatalytic domain. Also included are a nucleic acid comprising a gene, a ccatalytic domain have been the presence of an effector causes ct he nucleic acid to catalyse a reaction), isolating an RCANA (comprising a catalytic and an effector domain involving randomising at least one concleic acids that interact with the catalytic acids that interact with the catalytic target of the catalytic domain, adding an effector molecule to the nucleic acids and isolating ct those nucleic acids that interact with the catalytic target of the pacific modification and modifying the target using a RCANA, (involving providing a RCANA, capable of target cause a RCANA, specific activity), selecting an RCANA, and detecting an CCANA, (involving isolating an RCANA, capable of target cause a RCANA, especific activity), selecting an RCANA, and detecting an CCANA, (involving isolating an RCANA, capable of target cause a RCANA, specific activity), selecting an RCANA, and detecting an CCANA, (involving isolating an RCANA, capable of target catalytic activity of the nucleic acid upon exposure of the host cell to the effector. The RCANA is useful for regulate the expression of a reporter concleic acid upon exposure of the host cell to transcript to alter its coding potential, and a regulatory domain which comprises a catalytic domain, that modifies a tracell to result to the constituct into a host cell and measuring the constituct in the catalytic activity of the product. The concentration of the product in the constitution of the product is coding potential, and a regulatory domain which reports to alter its coding potential, and a regulatory domain which the catalytic domain of the effector is partially inhibited according to the concentration of the gene. The product is produced in a metabolic pathway that is being regulated, the product is chosen from protein, enzyme, protein pharmaceutical, metabolite, drug, dye, vitamin, food additive, chemical additive, pesticide, insecticide, feed compound, and a waste product. The drug (RCANA) segment comprising an effector domain and a nucleic acid catal domain in which one or more critical catalytic residues of the nucleic The invention relates to a regulatable, catalytically active nucleic acid acid catalyst

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RESULT 115
ADF12405
ADF12404
XX
AC ADF124
XX
DT 12-FEJ
XX L1 re
XXX
Gene
KW gene
KW retrat
XXX Homo
XXX Homo
XXX Homo
XXX Homo
XXX HOM
PN US20
XXX L5-
PR 15-
PR 15-
PR 28-
PR 28-
PR 01-
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PT 15-
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Best Local S
Matches 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Creating an insertional mutation in the germ line of an animal, useful for generating a mutation in an offspring of an animal, comprises introducing into an animal a nucleic acid molecule comprising a germ l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy; insertional mutation; germ line specific promoter;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-AUG-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mutation generation;
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96US-00749805.
97US-00847844.
2000US-00653812.
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g terminal repeats;
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human; primer;
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The invention describes a method of creating an insertional mutation in the germ line of an animal by introducing into an animal a nucleic acid molecule comprising a germ line specific promoter. The method is useful for generating a mutation in an offspring of an animal, or for isolating a nucleic acid from a genome of an offspring of an animal. The method may also be used to correct genetic defects in animals, especially humans. The nucleic acid is useful for generating mutations in a cell for assessing the frequency with which selected cells under go insertional mutagenesis for the generation of transgenic animals. This sequence represents a primer used to characterise the insertion site of the injection of the contraction of the contraction

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Example

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SEQ ID

NO 151; 102pp;

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RESULT 116
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The present sequence exxemplified the primer of the invention, and is used to sequence Incyte clone 1 (AN42737). The primer of the invention is labelled with a set of at least 2 different fluorescent labels. The set comprises an energy-transfer fluorescent label with at least 1 each of a donor fluorophore and an acceptor fluorophore capable of energy transfer, and separated by a distance x, and a second similar fluorescent label in which the separation distance is y, x and y being sufficiently
                                                                                                                                                                                                                      Set of energy-transfer fluorescent labels with donor and different separations - useful for DNA sequencing allows analysing wavelengths or an increased throughput.
                                                                                                                                                                                   Example 1;
                                                                                                                                                                                                                                                                                                                 WPI; 1998-414127/35.
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                                                                                                                                                                                   Page 14;
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Pred. No. 2.
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RESULT 117
AXX84259/c
ID AAX842
XX AAX842
XX OB-SEP
XX PCR PX
XX NCk as
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               different for the two fluorescent labels to produce distinct fluorescent signals. Fluorescent labels are useful in multicomponent analyses, e.g. as probes for fluorescent in situ hybridisation or especially as primers
                            AAT32790 standard;
                                                                                                                                                                                                            Sequence 25 BP; 1 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                    This sequence represents a PCR primer used to isolate DNA encoding the human Nck associated protein 1 (Nap1) of the invention. Nap1 inhibits apoptosis. The protein can be used in the investigation, diagnosis and treatment (e.g. by gene therapy) of Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                     Protein inhibiting apoptosis, useful in the diagnosis and treatment Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                         Sakaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nck associated protein 1; Napl; human; apoptosis; Alzheimer's disease; therapy; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                        Disclosure; Page 76; 90pp; Japanese.
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(SAKA/) SAKAKI Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA sequencing
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                            DNA; 26 BP
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Pred. No. 2.6e
0; Mismatches
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                                                                                                                                                                   Score 19.2; DB 1;
Pred. No. 2.6e+02;
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RESULT 119
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XC AAT32778;
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DT 18-FEB-19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Double-stranded (ds) DNA can be purified from complex mixtures of nucleic acids, proteins, endotoxins, nucleases, etc. by passing the mixture over a support to which an oligonucleotide is covalently attached; the oligonucleotide is solven to form a triple helix by hybridisation with a specific sequence present in the dsDNA. The method is particularly suited to purification of plasmid DNA. In an example, the present oligonucleotide was used for purifying plasmid px12726 (specially constructed by inserting a linker comprising a (GA) 25 homopurine sequence into the BamHI and ECORI sites of pBKS+)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Triple helix-forming
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-FEB-1997
                                                                                                                    purification;
                                                                                                                                Triple helix;
                                                                                                                                                          Triple helix-forming oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Purificn. of double stranded DNA by triple helix formation - on hybridising immobilised oligo-nucleotide to specific sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9618744-A2
                                         20-JUN-1996.
                                                                 WO9618744-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (RHON ) RHONE POULENC RORER SA
                                                                                                                                                                                                                                                                                                                       1180 AGAGAAAGAGAGAGAGAAATCA 1203
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                                                                                                                                                                                                                                                                                                         26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7; Page 18; 34pp; French.
                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                         AGAGAGAGAGAGAGAGAGAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Scherman
                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                    (first
                                                                                                                    triplex formation; Hoogsteen base pairing; plasmid; double-stranded DNA; homopyrimidine; polypurine; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            triplex formation; Hoogsteen base pairing; plasmid;
double-stranded DNA; homopyrimidine; polypurine; pXL2726;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94FR-00015162.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            95WO-FR001468
                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                     0.4%;
                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotide for purifying plasmid pXL2726
                                                                                                                                                                                                                                         26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wils P;
                                                                                                                                                                                                                                         BP.
                                                                                                                                                                                                                                                                                                                                                          Score 19.2; DI
Pred. No. 2.6e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                        2.6e+02;
3;
                                                                                                                                                                                                                                                                                                                                                                                       명
                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                         26;
                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                comprises
in target
                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                             0;
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08-NOV-1995;

95WO-FR001468

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RESULT 120
AAD03682/c
ID AAD036
XX AAD036
AC AAD036
XX Human
XX Human;
XW Human;
XW Antips
XW antips
XW Sperma
XW Sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EXEXEXE EFFX EXSSSSSSSSSSSSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Double-stranded (ds) DNA can be purified from complex mixtures of nucleic acids, proteins, endotoxins, nucleases, etc. by passing the mixture over a support to which an oligonucleotide is covalently attached; the oligonucleotide is able to form a triple helix by hybridisation with a specific sequence present in the dsDNA. The present sequence is a preferred oligonucleotide which can form a triple-helix with the homopurine target sequence (GA) 25. The target sequence may be present naturally, e.g. in a plasmid origin of replication, or can be introduced artificially. The method is particularly suited to purification of plasmid DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; phosphodiesterase; PDE; zcytor13; antiasthmatic; antiarthritic; antiporiatic; cytostatic; antiatherosclerotic; antiinfertility; cardiant; antiinflammatory; dermatological; wound healing; antiviral; antibacterial; therapy; inflammatory bowel disease; diverticulitis; spermatogenesis; sperm capacitation; immunocontraceptive; vaccine; cancer; reperfusion ischaemia; psoriasis; melanoma; myocarditis; PID;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pelvic inflammatory disease; eczema; scleroderma; vasoconstriction; heart arrhythmia; congestive heart disease; muscle spasm; fatigue; chromosomal abnormality; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 26 BP; 1 A; 11 C; 2 G; 12 T; 0 U; 0 Other;
   Novel human phosphodiesterase polypeptide, zcytor13 and polynucleotide
                                                                                                                                                                                                                                                                                                                   06-OCT-2000; 2000WO-US027734.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD03682
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD03682 standard; DNA; 26 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 13; Page 26; 34pp;
                                                                WPI; 2001-266312/27.
                                                                                                                                                                                                                                                     07-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                             12-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200125444-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-JUN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Purificn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1996-300660/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-DEC-1994;
                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (RHON ) RHONE POULENC RORER SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1180 AGAGAAAGAGAGAGAGAAATCA 1203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               full length zcytor13 cDNA isolating polyA PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21;
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                                                                                                                              SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGAGAGAGAGAGAGAGAGAGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Scherman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                     99US-00414025
                                                                                                                              Novak JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94FR-00015162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.4%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Á
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Wils P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 19.2; DB 1; Length 26; Pred. No. 2.6e+02; O; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZC7764b.
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e in target
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encoding it, for detecting human chromosomal abnormalities, identifying modulators and treating inflammatory and cardiovascular diseases
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1C; Page 118; 122pp; English.

cc and its corresponding protein. Zcytoril protein is used to promote wound come in the corresponding protein. Zcytoril protein is used to promote wound come in its corresponding protein. Zcytoril protein is used to promote wound come in its corresponding protein. Zcytoril protein is used to promote wound come in its corresponding protein and anti-viral effects and correct contents or antagonists or antagonists. Zcytoril its correct correct correct contents of inflammation, inflammatory heart correct correct contents. The protein muscle inflammation, inflammatory heart correct correc The patent discloses novel human phosphodiesterase (PDE), and its corresponding protein. Zcytor13 protein is used to zcytor13 cDNA

Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;

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S
문
                                     Matches
                                                      Query Match
                                            Local
                 5392 TAAAAAATACAAAAAGAAAAAA 5415
26 TAAAAAAAAAAAAAAAAAAA
                                              Similarity
                                     Conservative
                                            0.4%;
                                   Score 19.2; D
Pred. No. 2.6e
0; Mismatches
                                    0
                                              .6e+02
                                                      DB 1; Length 26;
                                     Indels
                                     0
                                     Gaps
                                     0
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RESULT 121
AAS20596/c
ID AAS205
XX AAS205
XX 23-APR
XX 24-Maman;
KW Human;
KW microb
KW gastro
KW acquir
KW acquir
KW acquir
KW acquir
XX 19-MAF
XX 17-MAF
XX 17-MAF
XX 27MO
XX Adler
XX Adler Human zsig63 cDNA sequencing primer 2C7764a. 23-APR-2002 AAS20596 AAS20596 standard; DNA; 26 (first entry) ₽₽

Human; zsig63; chromosome 4q12-4q13; salivary protein; antimicrobial; ss; microbial infection; tooth decay; periodontal disease; thrush; emphysema; gastrointestinal disease; urinary tract infection; vaginal infection; skin infection; epithelial wound; chronic tissue damage; cystic fibrosis; acquired immunodeficiency syndrome; AIDS; lung infection; sarcoidosis; chronic bronchitis; gene therapy; protein therapy; primer; ZC7764a.

Homo sapiens.

US6331413-B1

18-DEC-2001

17-MAR-2000; 2000US-00527345

17-MAR-1999; 99US-0124820P

(ZYMO) ZYMOGENETICS INC

Adler DA,

Sheppard

В Э

WPI; 2002-096707/13.

Polynucleotides encoding salivary proteins useful as anti-microbial

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RESULT 122
AAS1934/c
ID AAS193
XX AAS193
XX AAS193
XX Oligon
XX Synthe
XX Synthe
XX Synthe
YT repeat
FT re
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Best Local Similarity
Matches 21; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a polynucleotide derived from the 4q12-4q13 region of human chromosome 4 and encoding a zsig63 polypeptide, a secreted salivary protein with anti-microbial activity. Due to their microbial activity, the sequences can be used in the study of microbial infections, e.g. for recombinant production of anti-microbial proteins. The sequences can be used in the treatment of tooth decay, periodontal disease, thrush, gastrointestinal disease, urinary tract infections, vaginal infections, skin infections, epithelial wounds, chronic tissue damage, acquired immunodeficiency syndrome (AIDS), cystic fibrosis, lung infections, sarcoidosis, emphysema and chronic bronchitis. This sequence represents a sequencing primer for cDNA encoding human zsig63
                                                                                                                                                         Purifying double-stranded (ds) DNA from a solution containing dsDNA and other components, comprises passing the solution through a support comprising a covalently coupled oligonucleotide able to form a triple helix with the dsDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAS19344 standard;
                                                                                                          Example
                                                                                                                                                                                                                                                                                                                                                Crouzet J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        26-MAY-2000; 2000US-00580923
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAY-2001; 2001WO-US017122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                06-DEC-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAS19344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Col 53;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 repeat_region
                                                                                                                                                                                                                                                                                                                                                                                                    (AVET )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA purification;
                                                                                                                                                                                                                                                                                               2002-097772/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26 TAAAAAAAAAAAAAAAAAAA 3
                                                                                                             7.2;
                                                                                                                                                                                                                                                                                                                                                                                                       AVENTIS PHARMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TAAAAAAATACAAAAAAGAAAAAA 5415
                                                                                                                                                                                                                                                                                                                                             Scherman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                     Page 20; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers 5. .26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /rpt_type= TANDEM 5...6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence used to purify plasmid XL2726.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 triple helix;
                                                                                                                                                                                                                                                                                                                                                                                                       SA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             b
"CT repeat type"
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                                                                                                                                                                                                                                                                                                                                                   Wils P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ΒP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 19.2;
Pred. No. 2.
                                                                                                                                                                                                                                                                                                                                                Blanche F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                6e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 purification;
                                                                                                                                                                                                                                                                                                                                                   Cameron
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                        coupled oligonucleotide capable of forming a triple helix with the double-stranded DNA by hybridisation with a specific sequence present in the double-stranded DNA. The method is useful for purifying double-stranded DNA contained in a solution and mixed with other components. The new method is a simple, rapid and effective method for DNA purification, and makes it possible to obtain especially high purities with high yields. The method enables DNA to be purified from complex mixtures comprising other nucleic acids, proteins, endotoxins, nucleases and the like. The supports may be readily recycled, and the DNAs obtained display improved properties to pharmaceutical safety. Further, the method entails only one step contrary to prior art. The present sequence represents an of the contract of the method of the method of the contract of the contract of the present sequence represents an entitle contract of the method of the contract of the contract of the method of the method of the contract of the contract of the contract of the method of the method of the contract of the c
Sequence 26
                                                                                                                         invention
BP; 1 A;
11 C; 2
G; 12
Ή.
0 U;
0 Other;
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Watch 0.4%; Local Similarity 87.5%; Score 19.2; DB 1; Pred. No. 2.6e+02;

Query Match Best Local S Matches 21 21; Conservative 0; Mismatches Length 26; Indels 0 Gaps 0

밁 Ś

15-NOV-2002 ABS52638 standard; DNA; (first entry) 26 ВP

Human secreted salivary protein zsig63 PCR primer ZC7764a.

RESULT 123
ABS52638/c
ID ABS526
XX ALIMAN;
XX Human;
XX Human;
XX ALIMAN
XX Human;
XX Carinar
XX Carinar
XX US2002
XX US2002
XX US2002
XX ADILE;
X Human; secreted salivary protein; zsig63; immunogen; zsig63-cytokine; antibody-cytokine; in vivo killing; pathological microbe; bacteria; fungal; viral; infection; salivary gland; anti-microbial; dental carrie; cooth decay; periodontal disease; thrush; gastrointestinal disease; urinary tract infection; vaginal infection; skin infection; microfiora epithelial wound; pathogenic colonisation; invasion; pro-inflammatory; chronic tissue damage; vascular system; diabetes; anti-inflammatory; incompetent immune system; AIDS; acquired immunodeficiency syndrome; incompetent immune system; AIDS; acquired immunodeficiency syndrome; digestion; chemotherapy; radiation treatment; lung infection; cystic digestion; PCR; primer; ss. caries;

sapiens

US2002081701-A1

03-AUG-2001; 2001US-00922480.

17-MAR-1999; 17-MAR-2000; 2000US-00527345 99US-0124820P.

SHEP/) ADLER D A. SHEPPARD P O.

Adler DA, Sheppard 8

WPI; 2002-635468/68.

Novel secreted salivary protein, zsig63 and polynuseful for treating microbial infections, inflamma caries and lung infections associated with cystic zsig63 and polynucleotide encoding fections, inflammatory conditions, fibrosis.

Example 1; Page 29; 33pp; English.

The present invention relates to a new secreted salivary protein, zsig63 The invention is useful for detecting in a test sample, the presence of an antagonist or agonist of zsig63 protein activity. The invention is

This invention comprises a method of purifying double-stranded DNA from a solution containing the double-stranded DNA mixed with other components, comprising passing the solution through a support comprising a covalently

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RESULT 124
AAD45055/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC also useful as an immunogen for producing an antibody to zsig63 polypeptide. zsig63-cytokine fusion protein or antibody-cytokine fusion CC protein are useful for enhancing in vivo killing of target tissues. CC Pharmaceutical composition comprising purified zsig63 polypeptide are compressed purified zsig63 polypeptide are useful for treatment of dental caries (tooth decay), compressed purified and disease. The compressed purified protect periodontal disease, thrush and gastrointestinal disease. Other compressed purified and invasion. The invention is useful compressed in extremation and compressed purified a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; secreted salivary protein; zsig63 protein; host defense protein; immune modulating factor; antipathogenic; cell-cell signalling molecule; growth factor hormone activity, dental carries; infection; tooth decay; periodontal disease; gastrointestinal disease; infection; triush; urinary tract infection; vaginal infection; diabetes; obesity; thrush; urinary tract infection; vaginal infection; diabetes; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                anti-inflammatory; chronic tissue damage; lung dysfunction; restenosis; gene therapy; salivary gland dysfunction; prostate gland dysfunction; forensic DNA profiling; chondrosarcoma; atherosclerosis; primer; ss.
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WPI; 2002-642378/69
                                                                                                                                                                                                                                                                                                 17-MAR-1999; 99US-0124820P.
17-MAR-2000; 2000US-00527345.
                                                                                                                                                                                                                                                                                                                                                                                                                                03-AUG-2001; 2001US-00923236.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2002090677-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZC7764a primer used in the identification of human zsig63 DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 26 BP; 1 A; 0 C; 0
                                                                             Adler DA,
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                                                                                                                                                                   ADLER D A.
SHEPPARD P O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                             Sheppard PO;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.4%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
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Pred. No. 2.6e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .6e+02
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Novel secreted salivary polypeptide, zsig63, useful as antimicrobial agent for treating microbial infection, dental carries, periodontal disease, thrush gastrointestinal disease, and for aiding digestion.
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Example 1; Page 30; 33pp; English.

Cytokine, or as secreted extracellular matrix associated proteins with CC growth factor hormone activity. It is useful for treating conditions CC associated with pathological microbes, including bacterial, fungal and CC viral infections, for treating dental carries (tooth decay), periodontal CC disease, thrush and gastrointestinal disease, for treating urinary tract CC infection, vaginal infection and for preventing infection in skin and CC other epithelial wounds. zsig33 is useful for establishing normal CC microflora and protect against pathogenic colonisation and invasion, for treating chronic tissue damage e.g. damage in extremities associated with CC diabetes and useful as anti-inflammatory agents. It is useful as a marker CC of lung dysfunction, salivary gland dysfunction, or dysfunction of CC prostate gland. It is also therapeutically useful for aiding digestion. CC plynucleotides of the invention are used in gene therapy for increasing CC or inhibiting zsig3 activity, for detecting abnormalities on human CC diagnostics in forensic DNA profiling. Sequences of the invention are used in gene therapy for cardiac CC myocytes, for proliferation or differentiation of cardiac CC myocytes, for proliferation or differentiation of cardiac combibiting chondrosarcomas, atherosclerosis, restenosis and obesity. The human resident contraction of adaption of the human resident contraction of a cardiac conhibiting chondrosarcomas, atherosclerosis, restenosis and obesity. The can be used in detecting agoniets and antagoniets of its activity, and also useful as a host defense polypeptide, immune modulating factor, antipathogenic polypeptide, cell-cell signalling molecule, growth factor cytokine, or as secreted extracellular monecule, growth factor. present sequence is a primer used in the identification of human zsig63 invention relates to human secreted salivary polypeptide designated growth factor, 18

Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;

Query Match Best Local S Matches 21 21; Similarity Conservative 0.4%; 0; Mismatches Score 19.2; Pred. No. 2. .6e+02 DB 1; Length 26; Indels 0 Gaps 0

Ś 밁 5392 TAAAAAATACAAAAAAGAAAAA 5415 26 TAAAAAAAAAAAAAAAAAAA 3

RESULT 125
AAS20671/c
ID AAS206
XX AAS206
XX O9-APR
XX Cytoki
KW Inatura
KW Inatura
KW B-cell
KW Inmunc
XX Inmunc
XX O9-MAI
PR 09-MAI
PR 01-UII
XX O1-UII
XX O1-UII AAS20671 AAS20671 standard; DNA; 26 ₽P

Human zalphall Ligand sequencing primer 09-APR-2002 (first entry)

Cytokine; zalphall Ligand; zalphall receptor; NK cell progenitor; natural killer cell proliferation; T-cell proliferation; B-cell proliferation; anti-tumour response; immune system; B-cell proliferatimmunostimulant; cytostatic; human; sequencing primer; ss

Homo sapiens

US6307024-B1

23-OCT-2001

09-MAR-2000; 2000US-00522217.

09-MAR-1999; 11-MAR-1999; 01-JUL-1999; 99US-0123547P. 99US-0123904P. 99US-0142013P.

(ZYMO) ZYMOGENETICS INC

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to the isolation of a novel cytokine, CC zalphall Ligand and the polynuclectide encoding it. The invention also CC gives the sequence for the zalphall receptor and the polynucleotide CC encoding it. The zalphall Ligand polypeptide stimulates proliferation of Encoding it. The zalphall Ligand polypeptide stimulates proliferation of RC cells, proliferation of Freells, proliferation of B-cells etimulated with CC anti-CD40 antibodies, stimulates an antigenic response in a mammal, and CC reduces proliferation of B-cells stimulated with anti-IgM antibodies. The CC zalphall Ligand polypeptide is also useful in preparing antibodies that bind to zalphall Ligand epitopes. The zalphall Ligand polymucleotides can be used as probes or primers to clone regions of a zalphall Ligand gene, CC and in gene therapy. Zalphall Ligand may also be used to identify combitors of its activity, to enhance the generation of anti-tumour responses with or without the infusion of donor lymphocytes, and to continue the infusion of sevents required represents a continue regions of a salphall and the infusion of activate or stimulate the immune system. The present sequence represents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                            periodontal disease; thrush; gastrointestinal disease; epithelial wound; urinary tract infection; vaginal infection; skin infection; primer; pro-inflammatory; chronic tissue damage; vascular system; diabetes; AIDS; lung infection; cystic fibrosis; lung dysfunction; digestive; salivary gland carcinoma; Pneumocystis carinii infection; emphysema; chronic bronchitis; prostate dysfunction; prostate adenocarcinoma; cell culture media; gene therapy; human chromosome 4q12-4q13; dentinogenesis imperfecta; dentin dysplasia type II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novak
Gross
               17-MAR-1999; 99US-0124820P
17-MAR-2000; 2000US-00527345
                                                                       03-AUG-2001; 2001US-00922469
                                                                                                                                                                                                                                                                                                                                                               ss; PCR; zsig63; adhesin; salivary gland; dental carries;
                                                                                                                                                                                                                                                                                                                                                                                                     Human zsig63
                                                                                                                                                                                                                                                                                                                                                                                                                                         28-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABX93599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABX93599 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   a sequencing primer used to sequence CDNA clones in the isolation of human zalphall Ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 7; Col 139; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New zalphall ligand polypeptides and polynucleotides, useful for stimulating proliferation, activation, differentiation and/or induction of inhibition of specialized cell function, or for stimulating an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-040208/05
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Johnston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                     PCR/sequencing primer ZC7764a.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ĮĶ,
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Nelson AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19.2;
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Dillon SR, 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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RESULT 127
ADD33380
ID ADD333
XX
AC ADD333
XX
DT 15-JAN
XX
DT 15-JAN
XX
MOUSE
XX
MOUSE
KW ds; mc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an isolated zsig63 polypeptide comprising at CC least 90% identity to an amino acid sequence which comprises domain 1 of CC zsig63 domain 2, domain 3, mature zsig63 and full length zsig3. Also concluded are the polynuclectide encoding zsig63, a zsig63 expression CC included are the polynuclectide encoding zsig63, a zsig63 expression CC a DNA encoding a fusion protest in (comprising amino acids 1-15, 16-37, 38-CC zsig63 reporter gene construct to identify zsig63 agonists, and producing CC an anti-zsig63 antibody using zsig63 and an additional protest), using a CC zsig63 protein activity. Zsig63 has antimicrobial activity and since construct to identify zsig63 agonists, and producing CC anishits high expression in salivary gland, can be used for treating CC dental carries, periodontal disease, thrush, and gastrointestinal CC disease, urinary tract infections, vaginal infections as in infections and other epithelial wounds. The polypeptides can be used to establish CC normal microfilora and protect against pathogenic colonization and CC invasion. Zsig63 can also be used for providing pro-inflammatory activity for treating chronic, tissue damage particularly in areas having limited CC inflammocompromised AIDS patients or in individuals that have undergone CC chemotherapy, radiation treatment, for treating lung infections e.g. in CC yetic fibrosis. Detection of zsig63 polypeptide at relatively high CC marker of lung dysfunction. Zsig63 is also useful in diagnosing CC marker of lung dysfunction. Zsig63 is also useful in diagnosing CC marker of lung dysfunctions such as prostate conditions and may be used to replace serum that is commonly used in CC ulture media and may be used to replace serum that is commonly used in CC inhibit zsig63 and patients of detecting abnormalities on human CC culture. The DNA is useful in gene therapy applications to increase or inhibit zsig63 activity, and for detecting abnormalities on human CC culture media and useful served to isolate and sequence nucleic acdis
                                                                                                                                                                                                                                                                                                문
                                                                                                                                                                                                                                                                                                                                   S
                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 29; 32pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Adler DA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (ADLE/) ADLER D A.
(SHEP/) SHEPPARD P
                                                                                                                                                                                                                                                                                                                                       5392 ТАЛАЛАЛАТАСАЛАЛАЛАЛАЛАЛА 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2003-328428/31
                                                                                                                                                                                                                                                                                                26
                                                                                                                                                                                                                                                                                                                                                                            21;
                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human zsig63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sheppard PO;
                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 1 A; 0 C; 0
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                                                                                                                                                                                                                                                                                                                                                                                                  0.4%;
87.5%;
                                                                                                                                                                                                                                                                                                                                                                              Score 19.2; DB 1; Length 26; Pred. No. 2.6e+02; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 25 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 Other;
                                                                                                                                                                                                                                                                                                                                                                                  <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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ds; mouse; array; mitochondrial; mitochondrial disease; oxidative

hybridisation; energy-metabolism; phosphorylation dysfunction;

Mouse mitochondrial DNA sequence SEQ ID NO:1151.

15-JAN-2004

(first entry

ADD33380;

ADD33380 standard; DNA; 26 BP

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CC of the invention is useful for determining an expression profile of a conse or human sample containing nucleic acid, by contacting the array CC with the sample under conditions allowing selective hybridisation, and CC measuring hybridisation of nucleic acid in the sample to the array to CC produce an expression profile of a first labelled sample containing nucleic acid crelative to a second, differently labelled sample containing nucleic acid crelative to a second sample is a reference or a standard. An array is useful for determining an expression profile diagnostic of an energy-metabolism-cc related physiological condition. An array of the invention is useful for determining mitochondrial biology gene expression profiles of organisms, which are useful for determining expression profiles of diagnostic of energy metabolism-related physiological conditions, identifying biochemical conditions, which are useful for determining expression profiles of diagnostic of energy metabolism-related physiological conditions, can by stological conditions, identifying biochemical conditions, genes, and utations involved in such physiological conditions, can by stological conditions, and creating and/or monitoring the efficacy of cuseful for defining expression signatures or profiles for mitochondrial conditions, and creating and identifying animal models of human cenergy metabolism-related physiological conditions. An array is also cuseful for defining expression signatures or profiles for mitochondrial conditions. An array of the invention contains probes of genes containing and species that result from containing and species and spins. An array of the invention contains probes of genes contained to make the probes of the invention. Some sequences are not previously recognised to participate in mitochondrial biology. The sequences shown in ADD3224-ADD35260 represent murine mitochondrial DNA crosses and spins. An array of the invention. Some sequences are not profiles of the invention. Some sequences are not profiles of the
                                                                                       Query Match
Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel array comprising at least two isolated nuclectide molecules, each molecule having a sequence capable of uniquely hybridising to a nucleic acid molecule which is an expression product of a gene involved in mitochondrial biology. The array comprises two or more isolated nucleic acid molecules or spots, each molecule having a sequence chosen from sequence of 994 human probes and 2046 mouse probes. An array
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2; SEQ ID NO 1151; 201pp; English.
                                                                                                                                                                              Sequence 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   use in diagnosing pathologies and identifying biochemical pathways.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-300821/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-AUG-2001; 2001US-0316323P
31-AUG-2001; 2001CA-02356540
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-MAR-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYEM-) UNIV EMORY.
                                          1192 АGAGAGAAATCAGAGAAAAGGCAGG 1215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          containing probes for genes involved in mitochondrial biology, lear determining mitochondrial biology gene expression profiles
w
                                                                                                               Similarity
                                                                                                                                                                            BP;
                                                                                         Conservative
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                                                                                                          0.4%;
                                                                                                                                                                                 5 C; 8 G; 0 T; 0 U; 0 Other;
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                                                                                                               Score 19.2;
Pred. No. 2.
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                                                                                         Mismatches
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                                                                                                               2.6e+02
                                                                                                                                  DB 1; Length 26
                                                                                         Indels
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RESULT 128

Sequence 26 BP; 1 A; 0 C; 0 G; 25

T; 0 U; 0 Other

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ADH44608/c
ID ADH4460
XX ADH4461
XX ADH4461
XX Human;
XX Human;
XX Human;
XX Human;
XX Human;
XX Immunoux
XX Immuno

                                                                                                                                                                                                                                                                                        stimulating an immune response in a mammal exposed to an antigen or pathogen (comprising: (a) determining a level of antigen- or pathogen-specific antibody; (b) administering a composition comprising zalphall ligand polypeptide in a pharmaceutical vehicle; (c) determining a post administration level of the antigen- or pathogen-specific antibody; and (d) comparing the level of the antibody in (a) with (b), where an increase in the antibody level indicates stimulation of immune response). The method is useful for stimulating an immune response in a mammal exposed to an antigen or pathogen, and for enhancing anti-tumour activity resulting in a reduction in tumour progression, decrease in metastasis, or tumour stasis. The tumour may be a haematopoietic tumour, a lymphoma or a B cell tumour. The raphall ligand is useful for treating a wide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      exposed to an antigen or pathogen comprises administering a composition comprising mature zalphall ligand polypeptide comprising residues 32-162 of ADH44572 in a pharmaceutical vehicle. Also included are stimulating arimmune response in a mammal exposed to an antigen or pathogen (comprising: (a) determining (in)directly the level of antigen or pathogen present in the mammal; (b) administering a composition comprising zalphall ligand polypeptide in a pharmaceutical vehicle; (c) determining (in)directly the level of antigen or pathogen in the mammal; and (d) comparing the antigen or pathogen level in (a) with (b), where a change in the level indicates stimulation of immune response), and
                                                                                                           range of diseases arising from defects in the immune system, e.g. systemic lupus erythematosus, rheumatoid archritis, myssthemia gradiabetes, for boosting immunity to infectious diseases, treating immunocompromised patients, such as HIV+ patients and in improving vaccines. The present sequence is a sequencing primer used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-MAR-1999; 99US-0123547P.
11-MAR-1999; 99US-0123904P.
01-JUL-1999; 99US-0142013P.
09-MAR-2000; 2000US-00522217.
                                                                                exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 7; SEQ ID NO 38; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pathogen, useful for enhancing anti-tumor activity resulting in reduced tumor progression or metastasis, comprises administering zalphall ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Stimulating an immune response in a mammal exposed to an antigen or pathogen, useful for enhancing anti-tumor activity resulting in red
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-AUG-2001; 2001US-00923246
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             tumour progression; metastasis; tumour stasis; haematopoietic tumour; lymphoma; B cell tumour; systemic lupus erythematosus; rheumatoid arthritis; myasthenia gravis; diabetes; infectious disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; ss; Zalphall ligand; Zalphall receptor; immune response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human cDNA encoding Zalphall sequencing primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to stimulating an immune response in a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunocompromised patient, HIV infection; vaccine; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Presnell SR, Johnston JV,
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RESULT 129
ADIO0944/c
ID ADIO09
XX ADIO09
XX ADIO09
XX Zalpha
XX Zalpha
XX HIV; v
XX HOMO E
PN US2003
XX U
RESULT 130
ADP19767/c
ID ADP197
XX
AC ADP197
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AC ADP197
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DT 26-AUG
XX
DE Human
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Best Local Similarity
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Best Local Similarity
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Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a novel isolated zalphall ligand polypeptide. The polypeptide of the invention may be useful for boosting immunity to infectious diseases and treating immunocompromised patients, such as HIV patients, as well as in improving vaccines. The current sequence is that of the PCR primer which was used in the exemplification of the invention.
                                                                                                                                                                                              ADP19767 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
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HIV; vaccine; human; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequencing primer SEQ 38 used to analyse human zalphall ligand clone DNA.
                              Human zalphall ligand PCR primer seqid 38.
                                                                                    26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 7; SEQ ID NO 38; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New zalphall ligand polypeptides, useful for boosting immunity to infectious diseases, and treating immunocompromised patients, such as human immunodeficiency virus (HIV) patients, or in improving vaccines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-811003/76.
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Johnston JV,
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ilarity 87.5%;
Conservative
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Pred. No. 2.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 19.2; DB 1;
Pred. No. 2.6e+02;
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09-MAR-1999;
11-MAR-1999;
01-JUL-1999;
09-MAR-2000;
                                                                                                                                                                                  Novak JE,
Gross JA,
                                                                 New zalphall ligand polynucleotide and polypeptide molecules, useful for treating cancer, e.g. melanoma, solid tumor, hematopoietic tumor, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytostatic; zalpha11 ligand; pharmaceutical; cancer; immune response; melanoma; tumour; solid tumour; haematopoietic tumour; lymphoma; human;
                                                                                                                                       WPI; 2004-440401/41.
                                                                                                                                                                                                                                                                                                                                                                                                                        10-SEP-2003; 2003US-00659684.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2004110932-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
Example 7; SEQ ID NO 38; 111pp; English.
                                             Lympnoma.
                                                                                                                                                                                                                                                           (ZYMO ) ZYMOGENETICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer; 88.
                                                                                                                                                                                  Presnell SR,
Johnston JV,
                                                                                                                                                                                                                                                                                                         2000US-00522217
                                                                                                                                                                                                                                                                                                                          99US-0123547P.
99US-0123904P.
99US-0142013P.
                                                                                                                                                                                     Sprecher CA, Foster DC, Holly Nelson AJ, Dillon SR, Hammond
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The invention describes an isolated polypeptide comprising a sequence of CC amino acid residues that is at least 90 or 95% identical to residues 41 CC (Gln) to 148 (Ile), or 32 (Gln) to 148 (Ile) or 32 (Gln) to 148 (Ile) of a sequence of 162 amino CC acids (SEQ ID MO:2, human zalphall ligand), fully defined in the CC specification. Also described are: a pharmaceutical composition CC comprising the polypeptide, and a vehicle; a method of treating cancer in CC melanoma; a method of stimulating an immune response in a mammal with CC melanoma; a method of stimulating an immune response in a mammal bearing CC that encode amino acid residues cited above, where the polynucleotide comprising a sequence of nucleotides that binds a receptor comprising 538 amino acids, Cf fully defined in the specification; a pharmaceutically acceptable compositing the polynucleotide encoding, in a pharmaceutically acceptable compositing the polynucleotide encoding, in a pharmaceutically acceptable CC useful accounts a control element; and a DNA segment comprising the following operably linked CC least 10 nucleotides of the polynucleotide molecule comprising at CC cuseful for treating cancer, e.g. melanoma, solid tumour, haemacopoietic cumour, or lymphoma. This sequence represents a primer used in the expression cloning of human cytokine zalphall ligand.

SEXSEXE 밁 S RESULT 131 ADJ13038/c Query Match Best Local S Matches 21 ADJ13038 standard; Sequence 26 BP; 1 A; 0 C; 0 G; 25 T; 0 U; 0 Other; 5392 TAAAAAATACAAAAAGAAAAA 5415 26 21; Similarity TAAAAAAAAAAAAAAAAAA 3 Conservative DNA; 0.4%; 21 ; Score 19.2; D ; Pred. No. 2.6e 0; Mismatches .6e+02 DB 1; Length 0 Gaps 0;

Human DNA probe used to immobilise CpG methylated DNA SeqID 165

ADJ13038;

20-MAY-2004

(first entry)

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Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Analysis of chemical modification of DNA involves obtaining sample of to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                     Synthetic.
                                                                                                                            Sequence of scissile link probe MRC071 (HL).
                                                                                                                                                                                                 AAN70281;
                                                                                                                                                                                                                             AAN70281 standard; DNA; 27 BP
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 3
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                08-JUL-1987
                                         EP227976-A.
                                                                                              Hybridisation,
                                                                                                                                                      03-OCT-2002
26-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-874843/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-JUN-2001; 2001US-0301370P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-JUN-2002; 2002US-00184085.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US2003152950-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probe; ss; chemical modification; methylation; array; CpG island;
tumour suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  14-AUG-2003.
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                                                                                                                                                                                                                                                                                                                            2436 GGATGAGAAGGGGAGAGGT 2454
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MINNA J D.
LUBBKE K J.
BALOG R P.
                                                                                                                                                                                                                                                                                                   GGATGAGAAGGGGAGAGGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 165; 210pp; English.
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                                                                                                                                                     (revised)
(first entry)
                                                                                                probe; ss
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                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
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                                                                                                                                                                                                                                                                                                                                                        Score 19; DB
; Pred. No. 2.7
0; Mismatches
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RESULT 133
AANTO274/c
ID AANTO2
XX AANTO2
XX O3-OCT
DT 03-OCT
DT 26-MAY
XX Hybrid
XX Hybrid
XX Synthe
XX Synthe
PN EP2279
XX O8-JUL
XX O8-JUL
XX O1-DEC
XX O1-DEC
XX Duck F
XX D
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Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The patent claims a new molecule of formula (NA1---S---NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; --S--- a scissile linkage; n= 1 or 1,000, which is used for the detection of specific DNA or RNA sequences in a test soln. The scissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing OS field.)
  The patent claims a new molecule of formula (NA1---S---NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; ---S--- = a scissile
                                                                           Example; p29; 46pp; English.
                                                                                                                          Synthetic nucleic acid probes linked by a scissile linkage.
                                                                                                                                                                                                      WPI; 1987-186567/27.
                                                                                                                                                                                                                                                     Duck P,
                                                                                                                                                                                                                                                                                                                                                         05-DEC-1985;
                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-1986;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-OCT-2002
26-MAY-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic nucleic acid probes linked by a scissile linkage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1987-186567/27.
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                                                                                                                                                                                                                                                                                                       (MBIO-)
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                                                                                                                                                                                                                                                                                                       MEIOGENICS INC.
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                                                                                                                                                                                                                                                     Bender R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              scissile link probe
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(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probe; ss
                                                                                                                                                                                                                                                                                                                                                         85US-00805279
                                                                                                                                                                                                                                                                                                                                                                                                          86EP-00116906
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                                                                                                                                                                                                                                                        Стовьу
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
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Pred. No.
                                                                                                                                                                                                                                                           Robertson
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                                                                                                                                                       comprising two
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                                                                                                                                                       nucleic acid sequences
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RESULT 134
AAN92240/c
ID AAN922
XX AAN922
XX 25-MAR
DT 25-MAR
DT 25-APR
XX SPOOE
XX Probe
XX CPT Misc_f
FT Misc_f
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Matches 22
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                    Probe MRCO46 is bound by a permanent linkage to a solid support at its 3'end. It is used by reacting excess probe with a target nucleic acid, nicking hybridised probe at least once a predetermined sequence to form 2 or more probe fragments hybridised to the target sequence, which results in the probe fragments becoming hybridised to another probe; and identifying probe fragments, so detecting the target sequence. The probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SS probe MRCO46.
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31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAN92240 standard;
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                                                                                                                                                                                                                                                                                  Detecting target nucleic acid molecules - using excess complementary nucleic acid probes and nicking to complete a cycling sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-APR-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-APR-1988;
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                                                                                                                                                                                                                               Disclosure; Page 24; 34pp; English.
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(revised)
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17. .27
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Best Local S
Matches 22
Probe MRCO71 is bound by a hydrolysable linkage to a solid support at its 3' end. It is used by reacting excess probe with a target nucleic acid; nicking hybridised probe at least once within a predetermined sequence to form 2 or more probe fragments hybridised to the target sequence, which results in the probe fragments becoming hybridised to another probe; and identifying probe fragments, so detecting the target sequence. The probe can react with target sequence to complete a cycling sequence. Using this system, sensitivity of 10 exp. -19 to 10 exp. -20 molecules of target can be obtd. The probe is cleavable at the ribonucleotides by a ds RNase, eg RNase H or ExoIII. (Updated on 31-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 27 BP; 0 A; 0 C; 0 G; 21 T; 6
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31-OCT-2002
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18. .2
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RESULT 136
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RESULT 137
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Best Local Similarity
Matches 22; Conserv
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                           Synthetic
                                                                       Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothicate; ss.
                                                                                                                                                                                                                                                                      12-JUN-2001
                                                                                                                                                                                                                                                                                                                    AAF99706;
                                                                                                                                                                                                                                                                                                                                                                            AAF99706 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1993-171830/21
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Pred. No. 2.8e+02;
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                                                                                                                                                                                                                                         tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovalar glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; plaque neovascularisation; telangiectasia; haemophiliac joint; plaque neovascularisation; intestinal adhesion; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     against tumour antigens, viral antigens (e.g. herpesviridae, retrand/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The matching the staphylococcus, fungal antigens and/or parasitic antigens. The matching cancer, asthma, infectious disease, all also useful for preventing cancer, asthma, infectious disease, all
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequents one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 27 BP; 0 A; 0 C; 0 G; 27 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immune deficiency. The present sequence can also be used to redirect Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothicate backbone
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27-SEP-1999; 99US-0156135P
23-AUG-2000; 2000US-0227436P
                                                                                                 WO200253141-A2
                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                         scleroderma;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Angiogenesis inhibitory oligonucleotide #911.
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                                                                                                                                                                                                                         hypertrophic scar.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; phosphorothicate backbone; ss.
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                                                               (IOWA ) UNIV IOWA RES FOUND.
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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Pred. No.
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2.8e+02;
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Best Local S
Matches 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, oesophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, non-Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovarian cancer, pancreatic cancer, prostate cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for
                                                                                                                                                                                                                                                                                                                                                                                Human gene specific PCR primer #1103.
                                                                                                                                                                                                                                                                                                                                                                                                            02-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABK67015 standard; DNA; 27 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                              05-JAN-1999;
                                                                                                                                                                                                                                                                      05-MAR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                      ABK67015;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present sequence is an immunostir exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 developing cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell surface antigens
                                                                                                                                                                                                                   21-MAY-1997;
                                                                                                                                                                                                                                                                                                  US6352829-B1.
                                                                                                                                                                                                                                                                                                                                                       Primer; 88; DNA microarray; differential expression analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5389 ААТТАЛАЛАЛАТАСЛАЛАЛАСАЛАЛАЛ 5415
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                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
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                                                                                                                                                                                                                                              9905-00225928
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         312pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                immunostimulatory oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 2.86
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 19;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2.8e+02;
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The invention relates to producing a sub-population of labeled nucleic acids (NAs) comprising contacting a NA sample from a physiological source, with a pool of 50 distinct gene specific primers under suitable

differences in sources, using

Example 3; SEQ

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NO 1103; 11pp; English.

WPI; 2002-314699/35.

Chenchik A,

Jokhadze G,

Bibilashvilli

(CLON-)

CLONTECH

LAB INC.

Producing sub-population of labeled nucleic acids, useful for analyzing differences in RNA profiles between several different physiological sources, using set of distinct gene specific primers.

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RESULT 141
ACH03245/c
ID ACH032
XX ACH032
XX ACH032
XX Inmunc
XX Inmunc
XX Inmunc
XX Inflan
XX Inflan
XX US2003
XX US2003
XX 13-MAI
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PM 29-MAI
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PM (BERG,
XX WPI;
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Best Local (
                                                                                                                                                                                                        Treating non-allergic inflammatory diseases, such as psoriasis, ecz allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss
                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-521815/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAR-2001; 2001US-0279642P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAR-2002; 2002US-00112653
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACH03245 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BERG/)
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                                                                                                                                          Page 32; 229pp; English.
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Pred. No. 2.8e+02;
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The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory

RESULT 143 AAH28299/c

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RESULT 142
ADB37208/c
ID ADB372
XX ADB372
XX ADB372
XX Immuno
XX ds; al
KW ds; al
KW hypo-r
XX Synthe
XX US2003
XX US2003
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Best Local S
Matches 22
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Best Local 8
                                                                                                                                                                                                                                                  The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                                                                                                         Sequence 27
                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 17; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bratzler RL, Petersen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BRAT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ds; allergy; asthma; poly-G nucleic acid; aerosol formulation;
hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunostimulatory nucleic acid #822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (FOUR/)
                                              5389 AATTAAAAAATACAAAAAAGAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5389 AATTAAAAAAATACAAAAAAGAAAAA 5415
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27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ) BRATZLER R L.
) PETERSEN D M.
) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22;
                                                                                                                              Similarity
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AAAAAAAAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                      BP; 0 A;
                                                                                                Conservative
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                                                                                                                         0.3%;
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                                                                                                                                                                                                         0 C; 0 G; 27
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                                                                                                   0
                                                                                                                         Score 19;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fouron
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 19;
Pred. No.
                                                                                                   Mismatches
                                                                                                                                                                                                         T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ķ
                                                                                                                                                                                                            U; 0 Other;
                                                                                                                              2.8e+02;
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                                                                                                                                                       DB 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                   5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5
                                                                                                                                                  Length 27,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                                                                                                                      represents
                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                      0
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RESULT 144
AAH28297/c
ID AAH282
XX
AC AAH282
XX
DT 05-SEP
XX
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                                                                                                                                                                                                                                                                                                 The specification describes a method for partitioning endogenous cellular CC mRNA.protein (mRNP) complexes. The method comprises contacting a CC biological sample comprising mRNP complex with ligand that specifically binds a component of mRNP complex, separating mRNP complex by binding the CC ligand with a molecule specific for ligand, which is attached to the CC solid support and then collecting the mRNP complex by removing the CC complex from the support. The method is useful for in vivo partitioning CC of cellular mRNA protein complexes in a biological sample. The method is useful for determining the ribonomic profile of a cell which has numerous cuses including monitoring of tumour development, state of growth or state cf drug or toxin treatment and the state of cell aging or death, cC distinguishing ribonomic profiles among organisms, to discriminate complexes or toxin treatment and post-transcriptional contributions to gene expression and to track the movement of RNAs through RNP complexes, including the interactions of combinations of proteins with RNAs in RNP complexes. AAH28281-AAH28316 represent sequences derived from the 3' untranslated region (UTR) of RNAA of various genes. The sequences contain
                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 20
                    05-SEP-2001
                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 2 A; 1 C; 1 G; 0 T; 18 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Partitioning endogenous mRNA-protein complexes in vivo, by contacting sample comprising the complex with ligand that binds to a component of the complex and separating complex by binding ligand with a binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mRNA protein complex; tumour development; cell aging; death
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3' untranslated region sequence from neuronal cadherin gene.
                                                   AAH28297
                                                                                  AAH28297 standard;
                                                                                                                                                                                                                                                                                                                                untranslated region target sequences for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 6; Page 31; 49pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    motecule
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ribonomic profile;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                05-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAH28299 standard;
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                                                                                                                                                                                                   5394 AAAAAATACAAAAAAGAAAAAA 5415
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                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tenenbaum
                                                                                                                                                                                                                                   Conservative
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                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-0173338P
                                                                                                                                                                                                                                                                                                                                for RNA-binding proteins
                                                                                  RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA-binding protein; 88.
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                                                                                                                                                                                                                                                 0.3%;
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                                                                                  ВP
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                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                Score 18.8;
                                                                                                                                                                                                                                                 Pred.
                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                 No.
                                                                                                                                                                                                                                                 .9e+02
                                                                                                                                                                                                                                                                DB 1; Length 22;
                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                 Gaps
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3' untranslated region sequence
                                                                                                                                                                                            Partitioning endogenous mRNA-protein complexes in vivo, by contacting sample comprising the complex with ligand that binds to a component of the complex and separating complex by binding ligand with a binding
                                                                                                                                                                                                                                                                                          28-DEC-2000; 2000WO-US035583.
                                                                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                     mRNA protein complex; tumour development; cell aging; death; ribonomic profile; RNA-binding protein; ss.
                                                                                                                                                                                                                            WPI; 2001-425706/45
                                                                                                                                                                                                                                            Keene
                                                                                                                                                                                                                                                                          28-DEC-1999;
                                                                                                                                                                                                                                                                                                                        WO200148480-A1
                                                                                                                                                                                                                                                            (KEEN/) KEENE J D.
                                                                                                                                                                                                                                            à
                                                                                                                                                                                                                                            Tenenbaum
                                                                                                                                                                                                                                                                           99US-0173338P
                                                                                                                                                                                                                                            SA,
                                                                                                                                                                                                                                             Carson
                                                                                                                                                                                                                                                                                                                                                                             from neuronal cadherin gene
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of.

Example 6; Page 31; 49pp; English.

The specification describes a method for partitioning endogenous cellular CC mRNA-protein (mRNP) complexes. The method comprises contacting a CC biological sample comprising mRNP complex with ligand that specifically CC binds a component of mRNP complex, separating mRNP complex by binding the CI ligand with a molecule specific for ligand, which is attached to the CC solid support and then collecting the mRNP complex by removing the CC complex from the support. The method is useful for in vivo partitioning CC complex from the support end to the mRNP complex by removing the CC complex from the support. The method is useful for in vivo partitioning CC useful for determining the ribonomic profile of a cell which has numerous CC useful for determining the ribonomic profile of a cell which has numerous CC useful for determining the ribonomic profiles and system such as disease, CC drug or toxin treatment and the state of cell aging or death, CC distinguishing ribonomic profiles among organisms, to discriminate between transcriptional and post-transcriptional contributions to gene expression and to track the movement of RNAs through RNP complexes, including the interactions of combinations of proteins with RNAs in complexes. AAH28281-AAH28316 represent sequences derived from the 3' untranslated region (UTR) of mRNA of various genes. The sequences co sequences for RNA-binding proteins The sequences contain RNAs in RNP

Sequence 22 BP; 2 A; 1 C; 1 G; 0 T; 18 U; 0 Other;

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Query Match
Best Local Similarity
Matches 20; Conserv
 Conservative
        90.9%;
 0; Mismatches
          Score 18.8; DB 1;
Pred. No. 2.9e+02;
                    ۲.
                    Length
   Indels
  0
  Gaps
   0
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밁
              5394 AAAAAATACAAAAAAGAAAAA 5415
22
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RESULT 145
ADQ14522/c
ID ADQ145
XX ADQ145
XX ADQ145
XX ADQ145
XX Neuron
DE Neuron
XX Neuron
XX Metabo
KW metab
KW metab
KW metab ADQ14522 ADQ14522 standard; RNA; 22 ₽P

Neuronal-cadherin 3'-UTR consensus sequence SEQ ID NO:17.

23-SEP-2004

(first entry)

metabolic state; mRNA protein complex; mRNP complex; RNA binding protein; mRNA complex-associated protein; mRNP complex-associated protein; mRNA target; protein target; physiological pathway; neuronal-cadherin 3'-UTR consensus sequence; ss.

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RESULT 146
ADQ14524/c
ID ADQ145
XX ADQ145
XX ADQ145
XX ADQ16
AC AD
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method for assessing the metabolic state of a cell. The method comprises isolating at least one mRNP complex having at least one mRNP complex binding protein, and at least one mRNP or at least one mRNP complex-associated protein, and determining the expression level of the mRNP complex-associated protein, where the level of expression of the at least one mRNP complex-associated protein, where the level of expression of the at least one mRNP complex-associated protein is indicative of the metabolic state of the cell. The method can be used for assessing the metabolic state in a cell, and for identifying and evaluating mRNP and protein targets associated with mRNP complexes and implicated in the expression of proteins involved in common physiological pathways. The present sequence represents a neuronal-cacherin 3'-UTR consensus sequence, which is used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Assessing the metabolic state of a cell comprises isolating at mRNP complex comprising at least one RNA binding protein, and a one mRNA or at least one mRNP complex-associated protein.
                                                                                                                                                                                                                                                                                                                                                  metabolic state; mRNA protein complex; mRNP complex; RNA binding protein; mRNA complex-associated protein; mRNA complex-associated protein; mRNA target; protein target; physiological pathway; neuronal-cadherin 3'-UTR consensus sequence; ss.
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                                                                                               04-DEC-2003; 2003WO-US038475.
                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neuronal-cadherin 3'-UTR consensus sequence SEQ ID NO:19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-SEP-2004
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                                    04-DEC-2002; 2002US-00309788
                                                                                                                                                                                                                           WO2004057032-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5394 AAAAAATACAAAAAAGAAAAAA 5415
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Pred. No. 2.9e
0; Mismatches
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RESULT 147
ADQ145562/c
ID ADQ145
XX ADQ145
XX ADQ145
XX Neuron
XX Neuron
XX Metabo
KW MENA t
KW MENA t
KW MENA t
KW MO2004
XX WO2004
XX WO1; 2
XX ASSESS
PT MENP c
PT ONE MR
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Best Local &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               state of a cell. The method comprises isolating at least one mRNP complex having at least one mRNP complex sessociated protein, and at least one mRNP or at least one mRNP complex-associated protein, and determining the expression level of the mRNP complex-associated protein, where the level of expression of the at least one mRNP or mRNP complex-associated protein is indicative of the metabolic state of the cell. The method can be used for assessing the metabolic state in a cell, and for identifying and evaluating mRNA and protein targets associated with mRNP complexes and implicated in the expression of proteins involved in common physiological pathways. The present sequence represents a neuronal cadherin 3'-UTR consensus sequence, which is used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                               metabolic state; mRNA protein complex; mRNP complex; RNA binding protein; mRNA complex-associated protein; mRNP complex-associated protein; mRNP complex-associated protein; mRNA target; protein target; physiological pathway; neuronal-cadherin 3'-UTR consensus sequence; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Assessing the metabolic state of a cell comprises isolating at least mRNP complex comprising at least one RNA binding protein, and at least one mRNA or at least one mRNP complex-associated protein.
                                                                                                             WPI; 2004-525445/50.
                                                                                                                                                                                                                  04-DEC-2002; 2002US-00309788
                                                                                                                                                                                                                                                      04-DEC-2003; 2003WO-US038475
                                                                                                                                                                                                                                                                                         08-JUL-2004
                                                                                                                                                                                                                                                                                                                            WO2004057032-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neuronal-cadherin 3'-UTR consensus sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADQ14562 standard; RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22 BP; 2 A; 1 C; 1 G; 0 T; 18 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 4; SEQ ID NO 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Keene JD,
                                                                                                                                            Keene JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention describes a method for assessing the metabolic
                                                                                                                                                                                 (RIBO-) RIBONOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (RIBO-) RIBONOMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5394 ААЛАЛАТАСАЛАЛАДАЛАЛАЛ 5415
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAAAAATACAGAAAATAAAAA 1
                                                                                                                                            Tenenbaum
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                                                                                                                                            SA,
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                                                                                                                                                Carson
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Pred. No. 2.
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                                                                                                                                                Phelps
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least
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Assessing the metabolic state of a cell comprises isolating at least one mRNP complex comprising at least one RNA binding protein, and at least one mRNA or at least one mRNP complex-associated protein.

Example 4; Page 35; 86pp; English.

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RESULT 148
ADQ14560/c
ID ADQ145
XX ADQ145
XX ADQ145
XX ADQ145
XX Neuron
XX Neuron
XX MRNA C
KW MRNA C
PF 04-DEC
XX G
PF 04-DEC
XX (RIBO-
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Best Local (
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state of a cell. The method comprises isolating at least one mRNP complex having at least one mRNP complex thaving at least one mRNP complex expression at least one mRNP complex associated protein, and determining the expression level of the mRNP or mRNP complex associated protein, where the level of expression of the at least one mRNP or the at least one mRNP complex associated protein is indicative of the metabolic state of the cell. The method can be used for assossing the metabolic state in a cell, and for identifying and evaluating mRNP and protein targets associated with mRNP complexes and implicated in the expression of proteins involved in common
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  metabolic state; mRNA protein complex; mRNP complex; RNA binding protein; mRNA complex-associated protein; mRNA complex-associated protein; mRNA complex-associated protein; mRNA target; protein target; physiological pathway; neuronal-cadherin 3'-UTR consensus sequence; ss.
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                                                                                                                                                                                                                                                                                                 Example 4; Page 35; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2004-525445/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-DEC-2003; 2003WO-US038475
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADQ14560
                                                                                                                                                                                                                                                The present invention describes a method for assessing the metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RIBO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Tenenbaum SA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 2 A; 1 C; 1 G; 0 T; 18 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Carson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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Pred. No. 2.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Phelps
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Query Match Best Local S Matches 20

Similarity

90.3%;

Conservative

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Mismatches

2

Indels

0

Gaps

0;

Pred

Score 18.8; Pred. No. 3e G; 0 T; 0

3; DB 1; Length 3e+02;

26;

Sequence

26 BP;

24

A; 2 C; 0

Ç,

0 Other;

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RESULT 149
AAV12482
ID AAV124
XX AAV124
XX AV124
XX Synthe
XW Synthe
XW Synthe
XW Synthe
XW Synthe
PM US5714
XX 33-FEE
XX 15-APE
XX 15-APE
XX 15-APE
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XX CUYRP
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                                                   The present sequence represents an oligonucleotide used in an example of CC the present invention. The present invention describes a method for CC synthesising a selected oligonucleotide (I) having well defined ends. The CC method comprises: (a) annealing a primer to a single-stranded (ss) CC circular template to yield a primed circular template, where the template CC comprises: (i) at least one nucleotide sequence complementary to (I); and CC (ii) at least one nucleotide effective to produce a cleavage site in the CC cligonucleotide multimer; (b) combining the primed circular template with CC cligonucleotide multimer; (b) combining the primed circular template with CC cligonucleotide addition of auxiliary proteins to yield a so oligonucleotide comprising multiple copies of (I); and (c) cleaving the planucleotide comprising multiple copies of (I); and (c) cleaving the planucleotide comprising multiple copies of (I); and (c) cleaving the planucleotide comprising multiple copies of (I); and (c) cleaving the planucleotide comprising multiple copies of (I); and (c) cleaving the planucleotide ends. CC multimer at the cleavage site to produce (I) having well defined ends. CC for use, e.g. as probes and diagnostic agents and/or therapeutic agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5
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Best Local Similarity
Matches 20; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rolling circle synthesis of oligo:nucleotide(s) - using primed template to produce oligonucleotide multimer for cleavage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US5714320-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Col 45; 38pp; English.
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           В
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                           an example of the method of the invention for synthesising an RNA oligonuclectide, comprising combining a single-stranded circular oligonuclectide, comprising at least one copy of a nuclectide sequence complementary to the sequence of the desired RNA oligonuclectide with at least 2 types of ribonuclectide triphosphate and a polymerase enzyme to yield a single-stranded RNA oligonuclectide multimer complementary to the circular oligonuclectide the RNA oligonuclectide multimer comprises multiple copies of the desired RNA oligonuclectide. The methods can be used for producing RNA oligonuclectides having a specific sequence and well defined ends. The RNA oligonuclectides having a specific sequence and well defined ends. The RNA oligonuclectides produced can be used for modifying the structure or function of a target molecule. They can also be used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Page 36; 100pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthesis of oligo:nucleotide(s) - using a single-stranded circular oligo:nucleotide template ribonucleotide tri:phosphate(s) and a polymerase to form multimer(s) which can be cleaved.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1998-481202/41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            14-DEC-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV59215 standard; DNA; 26 BP
                                                                                                                                                                                                                                           Sequence 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The circular template was used for the synthesis of the oligomer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-FEB-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              therapeutic agent.
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                                                                                                                                                                                                                                                                                                        cleave disease-associated RNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYRP ) UNIV ROCHESTER.
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                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                              BP;
                                                                                                                       Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US003784
                                                                                                                                                                                                                                              24 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "Position 26 optionally bound to position 1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /"tag= a
/note= "Position 1 optionally bound to position 26"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag=
                                                                                                                                                    90.9%;
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                                                                                                                       0; Mismatches
                                                                                                                                                    Score 18.8; DB 1; Length Pred. No. 3e+02;
                                                                                                                                                                                                                                                                                                           DNA or protein
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                                                                                                                             Indels
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ADC65872
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AAX30018
IID AAX3000
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a detectably labelled oligonucleotide multimer, comprising multiple contiguous copies of a repeated oligonucleotides. The detectably labelled oligonucleotide multimer is useful for detecting a target molecule. Oligonucleotide multimers may be produced in sufficient quantity to be useful for diagnostic and medical applications. The multimers are useful for affinity labelling of proteins, and for signal amplification in highly sensitive affinity capture and sequence identification applications. The method provides a faster, cheaper and simpler way for large-scale production of DNA and RNA oligoners and multimers. The incorporation of labels enables the oligonucleotide multimers to be useful in diagnostics and medicine. The
RNA oligonucleotide synthesis; ribonucleotide triphosphate; polymerase; electroporation; calcium phosphate treatment; lipid-mediated delivery; cation-mediated delivery, bacterial infection; viral infection; drug resistant infection; double stranded DNA oligomer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New detectably labelled oligonucleotide multimer, comprising multiple contiguous copies of a repeated oligonucleotide - useful for detecting target molecules in diagnosis and medicinal applications.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Multimer; probe; diagnosis; synthesis; detection; polymerase; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX30018 standard; DNA; 26
                                                                                                                                                              DNA oligonucleotide
                                                                                                                                                                                                                               18-DEC-2003
                                                                                                                                                                                                                                                                                           ADC65872
                                                                                                                                                                                                                                                                                                                                                        ADC65872 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 26 BP; 24 A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   present sequence represents an oligonucleotide used in an example from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Page 41; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Precircle DNA oligonucleotide SEQ ID NO:5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the present invention
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                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.3%; Score 18.8;
90.9%; Pred. No. 3e
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RESULT 153
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                              synthesising an RNA oligonucleotide. The method is useful for oligonucleotide is introduced into the cell using direct injection, electroporation, calcium phosphate treatment, lipid-mediated delivery, or cation-mediated delivery. The method is useful for treating bacterial and/or viral infections in mammals, particularly drug resistant infections, and for producing double stranded DNA oligoners. The method is performed in the absence of an oligonucleotide primer, or without the addition of auxiliary proteins. This sequence represents an oligonucleotide used in the method of the invention.
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26-FEB-1997;
11-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method for synthesising an RNA oligonucleotide, comprising combining a single-stranded circular oligonucleotide template with at least two types of ribonucleotide triphosphate and a polymerase enzyme to yield a single-stranded RNA oligonucleotide multimer complementary to the circular oligonucleotide template, where the RNA oligonucleotide multimer comprises multiple copies of the desired RNA oligonucleotide. The method is useful for
                                                                    Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                        ABN13299 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthesizing RNA oligonucleotide involves combining single-stranded circular oligonucleotide, ribonucleotide triphosphate and polymerase enzyme to yield desired RNA complementary to circular oligonucleotide
                                            Homo sapiens.
                                                                                                                              Human GDMLP-1 25-mer scanning SEQ ID NO:5 sequence SEQ ID NO:13291.
                                                                                                                                                              29-MAY-2002
                                                                                                                                                                                           ABN13299;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; SEQ ID NO 5; 78pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2003087241-A1
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                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                 26
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95US-00393439.
97US-00805631.
2000US-00569344.
                                                                                                                                                             (first
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                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                A; 2 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                   0.3%;
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                                                                                                                                                                                                                                                                                                                                                                     Score 18.8;
Pred. No. 36
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                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 26;
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                                                                                       heart disease
                                                                                                    hGDMLP-1; heart;
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RESULT 154
ABV80872
ID ABV808
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AC ABV808
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DT 03-JAN
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ABV80872;

ABV80872

standard;

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03-JAN-2003

(first entry)

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                                                                                                                                                                                                                                                            The present invention describes a human genome-derived myosin-like CC protein 1 (hGpMLP-1). The protein and polynucleotide sequences of hGDMLP-CC 1 can be used in gene therapy and vaccine production. The hGDMLP-CC nucleic acids can be used as probes to detect, characterise and quantify CC hGDMLP-1 nucleic acids in samples, as amplification substrates, to CC provide initial substrates for the recombinant engineering of hGDMLP-1 CC protein variants having desired phenotypic improvements, and for CC expressing the proteins. The hGDMLP-1 proteins or polypeptides may be CC used as immunogens to raise antibodies that specifically recognise hGDMLP cc used as immunogens to raise antibodies that specific biomolecule CC and/or amount specifically of hGDMLP successing the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule CC capture probes for surface-enhanced laser desorption ionisation, as CC therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The CC polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a CC and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.
                                                                 Query Match
Best Local Similarity
Matches 21; Conserv
                                                                                                                                        Sequence 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-MAY-2000;
21-SEP-2000;
                                                                                                                                                                                            The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 13291; 214pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 desorption ionization,
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30-JAN-2001;
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30-JAN-2001;
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           w polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, as specific biomolecule capture probes for surface-enhanced laser sorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                          ftp.wipo.int/pub/published_pct_sequence
                                 3475
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2002-179446/23.
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                AGCAGACGGAAACCAAGTGTGATGA 3499
AGCAGAGTGAAGCCAAGTGTGAGGA 25
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2000US-0236359P.
2000US-0236359P.
# 2000UG-0236359P.
# 2001WO-US000661.
2001WO-US000662.
# 2001WO-US000664.
# 2001WO-US000665.
# 2001WO-US000665.
# 2001WO-US000665.
# 2001WO-US000666.
# 2001WO-US000669.
                                                                     Conservative
                                                                                                                                          BP; 9 A; 3 C; 10 G; 3 T; 0 U; 0 Other;
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                                                                                  0.3%;
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                                                                   0; Mismatches
                                                                                    Score 18.6;
Pred. No. 3.
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RESULT 155
ABS75627
ID ABS756
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AC ABS756
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                                                                                                                                                                                                                                                                                                                 The present invention relates to human testis expressed Patched like CC protein (HTPL, see ABV876759 to ABP78762 and ABB98519 to ABB98520). HTPL CABB98520 two isoforms, with a few single base pair differences between the CC two. One of the single base pair changes introduces a premature stop CC codon in HTPL-S (S for short) compared to HTPL-L (L for long). HTPL CC shares an overall structure organisation with the Patched protein. The CC shared structural features strongly imply that HTPL plays a role similar to that of Patched, and is a potential tumour suppressor. HTPL is CC important in regulating male germ cell development, and the HTPL gene was CC important in regulating male germ cell development, and the HTPL gene was CC important in regulating a disorder caused by mutation in HTPL, and in CC therapy and manufacture of a medicament for treatment or prevention of CC such disorder associated with decreased expression or activity of human CC HTPL. Such disorders include disorders of testis, or adrenal, adult and CC foetal liver, bone marrow, brain, kidney, lung, placenta, prostate, compared to human cation. HTPL proteins and nucleic acids are clinically useful diagnostic markers and potential therapeutic agents for committee the invention. The present oligonucleotide was used in an analysis of the strility and cancer. The present oligonucleotide was used in an accomplishment of the strility and cancer. The present oligonucleotide was used in an accomplishment of the strility and cancer.
                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 21; Conserv
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   ABS75627;
                                    ABS75627 standard; DNA; 25 BP
                                                                                                                                                                                                                                                                      Sequence
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30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AEOM-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 el isolated human testis expressed Patched like protein (HTPL), use identifying agonist and antagonist and specific binding partners, treating subjects having defects in HTPL.
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                                                                                                                                                             770 GCGCCCAAGCCCAGGAAGGGGCAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2; Page 341; 718pp; English.
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nilarity 84.0%;
Conservative
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2001WO-US000665.
2001WO-US000668.
2001WO-US000668.
2001WO-US000669.
2001US-0327898P.
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                                                                                                                                                                                                                                                                                                       invention
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Pred. No. 3.2e
0; Mismatches
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                                                                                                                                                                                                                                                                      0 Other;
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                                                                                                                                                                                                                                 1; Length
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RESULT 156
ACC45214/c
ID ACC452
XX ACC452
XX ACC452
XX 16-JUN
DT 16-JUN
XX Human
XX Homon
XX Homon
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Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel isolated nucleic acid that encodes one of three new isoforms of human pregnancy associated plasma protein E, PAPP-E. The products of the invention have abortive and contraceptive activity and can be used for gene therapy or in a vaccine. The nucleic acid, polypeptide encoded by it, or antibody to the polypeptide can be used in pharmaceutical compositions or vaccines for preventing or aborting pregnancy. PAPP-E is used in the antenatal diagnosis of dysgenetic pregnancies. The nucleic acids are used as probes to assess the level of PAPP-E isoform mRNA in chorionic villus samples, and the antibodies can be used to assess the expression levels of PAPP-E isoform proteins in chorionic villus samples, to diagnose dysgenetic pregnancies antenatally. This sequence represents an oligomer used in scanning the human PAPP-E genes described in the disclosure of the invention
                                                Human; cytostatic; nootropic; neuroprotective; antiinflammatory; antisense therapy; NAC; DBFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                 Human NAC chimeric phosphorothioate oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated nucleic acid encoding an isoform of human pregnancy associated plasma protein E, for preventing or aborting pregnancy
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                                                                                                                                                                     16-JUN-2003
                                                                                                                                                                                                                                     ACC45214 standard;
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(SHAN/) SHANNON M
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                                                                                                                                                                                                                                                                                                                                                                                    21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      man; pregnancy associated plasma protein B; abortive;
ive; gene therapy; vaccine; pregnancy; antenatal; dia-
pregnancy; primer; ss.
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                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                       DNA;
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Pred. No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Other;
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                                                                                                                                      NO: 74
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RESULT 157
AAQ20032/c
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                                                                                                                                                                                                                      The present invention describes a compound (I) 8-50 nucleobases in length CC targeted to a nucleic acid molecule encoding NAC, where the compound compound the profit of the specifically hybridises with the nucleic acid molecule encoding NAC and compound specifically hybridises with the compound specifically hybridises with the teast an 8-nucleobase portion of an active site on a nucleic acid compound composition comprising (I) composition comprising (I) and a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC (I); and (3) treating an animal having a disease or condition associated with NAC comprising administering (I) to the animal so that expression of CC (I); and (3) treating an animal having a disease or condition associated cC with NAC compounds (I) has cytostatic, nootropic, neuroprotective and CC antisense compounds (I) are useful for modulating the expression of NAC, e.g. hyperproliferative disease, neurological disease, or a disease (CC and for treating a disease or condition associated with expression of CC (NAC, e.g. hyperproliferative disease, neurological disease, or a disease (CC and sorder arising from abservant apoptosis. The compounds are also CC useful as research reagents and kits, or for diagnostics, therapeutics and prophylaxis, e.g. to prevent or delay infection, inflament-forming CC (CDA-like apoptosis protein (DEFCAP). NAC is also known as a death effector filament-forming content of the present sequence represents a human NAC chimeric or heapfurthicate antisense oligonuplatide which is civen in the
                                                                                                                                              Matches
                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                    Sequence 20
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                                                                                                                                                                                                                                                         phosphorothicate antisense oligonuclectide, which exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ISIS-) ISIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antisense compounds, useful for modulating the expression of NAC treating a disease or condition associated with the expression of e.g. hyperproliferative disease or neurological disease.
                                                                                                          3466 CTCATCTTCAGCAGACGGAA 3485
                                                                        20
                                                                                                                                                Conservative
                                                                                                                                                                                                                    BP; 5 A; 4 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "
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/mod_ba
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1. .5
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'note= "2'-O-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                                             95.0%;
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e= "2'-O-methoxyethyl (2'-MOE) gapmer"
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                                                                                                                                                               Score 18.4;
Pred. No. 3
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modified_base
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aziridinylcytosine; cross-linking group; tumour necrosis factor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cross-linking oligomer 212 for targetting human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ20032 standard; DNA; 22 BP
                  New sequence-specific non-photo-activated crosslinking agents - bind to the major groove of duplex DNA and are esp. useful for treating latent infections e.g. HIV.
                                                                Matteucci
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14-JAN-1991;
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                                                               Krawczyk
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91US-00640654.
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                                                                                                                                                                                                                                                                                                                                                                                                         "N4N4-ethanocytosine"
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Example 4; Page 25; 42pp; English

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RESULT 158
AAQ30380/c
ID AAQ303
XX AAQ303
XX AAQ303
XX Oligom
XX Oligom
XX Tumoux
XX Tumoux
XX Synthe
XX Key
FT modifi
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Matches 19
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07-DEC-1992
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 11-JUN-1992.
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/mod_base= OTHER
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== "OTHER= N6 methyl-8-oxo-2'
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0; Mismatches
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RESULT 159
AAQ30381/c
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XX AAQ303
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XX 25-MAR
DT 07-DEC
XX Oligom
XX Tumour
KW Tumour
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08-APR-1991;
17-APR-1991;
17-APR-1991;
17-APR-1991;
27-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human tumour necrosis factor beginning at nucleotide 1137 contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV; HER; HIV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                       Tumour necrosis factor; herpes simplex; AIDS; modified; HIV; RSV; HPV, malignancy; hepatitis; inflammation; ss.
                                                                                                                                                                                                                             Oligomer TNF212 for forming triplex with HUMINFAA target duplex.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New oligomers contg. modified bases -doublet in a DNA duplex, for treating herpes malignancy and inflammation.
            modified_base
                                                               modified_base
                                                                                                                     modified_base
                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                         25-MAR-2003
07-DEC-1992
                                                                                                                                                                                                                                                                                                                          AAQ30381 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1992-217083/26.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-NOV-1990;
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91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
91US-00766733.
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95.0%;
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Pred. No. 3.
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                                                                               N4
                          methyl-8-oxo-2' deoxyadenine"
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                                                                               ethanocytosine"
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Matches
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17-APR-1991;
17-APR-1991;
                                                                                                                                                            The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human tumour necrosis factor beginning at nucleotide 1137 contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV; HER; HIV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. See also AAQ2452-25501 and AAQ30226-448.
                                                                                                                                                                                                                                                                                                                                                                        Claim 12;
                                                                                                                                                                                                                                                                                                                                                                                                       New oligomers contg. modified bases -doublet in a DNA duplex, for treating herpes malignancy and inflammation.
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08-APR-1991;
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                                                                                                             Sequence 22
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                                                                                                                                                specimen to harsh conditions. See also AAQ254 (Updated on 25-MAR-2003 to correct PN field.)
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1182 AGAAAGAGAGAGAGAAAT 1201
                                    l Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                      Page 70; 77pp; English.
                                                                                                             BP; 3
                                    Conservative
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91US-00643382.
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/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
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3-00686546.
3-00686547.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Φ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matteucci MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          m5c
                                  0;
                                Score 18.4; DB 1; Length 22; Pred. No. 3.3e+02; O; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    N6 methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                            which form a triplex with G-C and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Milligan
                                    0
                                    Gaps
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5392 TAAAAAAATACAAAAAAGAAAAAA 5415

Query Match Best Local S Matches 20

. Similarity 20; Conserv

0.3%; nilarity 83.3%; Conservative

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Score 18.4; DB 1; Pred. No. 3.4e+02; Mismatches

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Length Indels

24;

0

Gaps

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RESULT 160
ABK48140/c
ID ABK4811
XX ABK481
XX ABR481
XX ABPERG
XX Aminog
KW Aminog
KW baked
XX Synthe
PN W02002
PD 28-FEE
XX Synthe
PN W22-AU
XX Synthe
PN W22-AU
XX Synthe
PN W23-AU
XX Synthe
PN W21-1
XX Examp
XX An is
PT compoi
XX An is
PT compoi
XX An is
PT compoi
XX Examp
CC the g
CC recom
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CC the g
CC hydro
CC the f
CC hydro
CC f hydro
CC Aspert
CC Aspert
CC Sever
                                                                The invention relates to an isolated polypeptide with aminopeptidase constructy and the gene encoding it (including sequences complimentary to the gene and which hybridise to it at high stringency), from Aspergillus CC niger. Also included are a nucleic acid construct comprising the above cc polynucleotide operably linked to one or more control sequences that CC direct the production of the polypeptide in a suitable expression host, a crecombinant expression vector comprising the above mucleic acid construct, a recombinant host cell comprising the above construct or corrector, and producing the protein comprising cultivating an above strain/ crecombinant host cell to produce a supernatant and/or cells comprising the polypeptide. The aminopeptidase is used to prepare a food composition such as dough to enhance the flavour of a baked product from the dough and for preparing a cheese to enhance the flavour. The invention provides a bacterial enzyme for protein comprising an above the flavour of a baked product from the dough and for preparing a cheese to enhance the flavour. The invention provides a bacterial enzyme for protein comprising the enzyme has been isolated and characterised, compared to a gents, and the enzyme has been isolated and characterised compared to a crevitually observed weak aminopeptidase activity, to produce flavouring constructions of enzymes to produce flavouring agents from constitute of the saminopeptidase activity which was detected in an entire of the saminopeptidase activity which was detected in an entire of the saminopeptidase activity which was detected in an entire of the saminopeptidase activity which was detected in an entire of the saminopeptidase activity which was detected in an entire of the saminopeptidase activity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABK48140;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK48140 standard;
  Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      An isolated polypeptide with aminopeptidase activity, compositions, such as bread and cheese, with enhanced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-257917/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aminopeptidase; primer; ss; food composition; dough; flavour enhancer; baked product; cheese; poly-T; reverse transcriptase PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Aspergillus niger aminopeptidase RT-PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-JUN-2002
                                                        structure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 40; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Basten D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (STAM ) DSM NV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-AUG-2000; 2000EP-00202995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-AUG-2001; 2001WO-EP009925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200216618-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AGAAAGAGAGAGAGAATT
                                                        of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dekker PJT,
     BP; 0 A; 0 C; 0 G; 23 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                      aminopeptidase gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Schuurhuizen PW, Schaap PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer poly-T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for preparing food flavoring.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Visser
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RESULT 161
ADM92894
ADM92894
AC ADM928
XX ADM928
XX SNP-Q
XX SNP-Q
XX SNP-Q
XX SNP-Q
XX SNP-Q
XX Homo

RESULT 162
AAS20595/c
ID AAS205
XX
AC AAS205
XX
DT 23-APR
XX
DB Human
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates an isolated polynucleotide (I) encoded by a cardiovascular associated (CA) gene, having allelic variation contained in a functional surrounding like full length cDNA for CA gene polypeptide, and with or without the CA gene promoter sequence. (I) is a polynucleotide comprising single nucleotide polymorphisms predicting cardiovascular disease. The polynucleotides are useful for assessing predisposition or susceptibility to a cardiovascular disease, e.g. atheroselerosis, ischemia/reperfusion, hypertension, restenosis, arterial inflammation, myocardial infarction, and stroke. These may also be used to predict personal medication schemes omitting adverse drug reactions, or as probes for detecting genetic polymorphisms and as templates for the recombinant production of normal or variant peptides/polypeptides encoded by the genes. This sequence corresponds to a PCR primer to amplify one of the genes of the invention.
   Human zsig63 cDNA sequencing primer ZC7231.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 75; 187pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New isolated polynucleotides comprising single nucleotide polymorphisms of the cardiovascular gene, useful for assessing predisposition or susceptibility to a cardiovascular disease, e.g. atherosclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JAN-2003; 2003WO-EP000060
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                                                                23-APR-2002
                                                                                                                                 AAS20595
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JAN-2002; 2002EP-00000153
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP-containing cardiovascular associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-JUN-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADM92894 standard; DNA; 25
                                                                                                                                                                                              AAS20595 standard;
                                                                                                                                                                                                                                                                                                                                                                                    1181 GAGAAAGAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2003-577532/54.
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                                                                                                                                                                                                                                                                                                                                                   μ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                   GAGAAAGAGAGAGAGAGA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   or stroke.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Schwers S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   13 A; 0 C; 12 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
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                                                                                                                                                                                              26
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                                                                                                                                                                                              먥
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 18.4; DB 1; Length 25; Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     #225.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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RESULT 163
ABS22637/c
ID ABS526
XX ABS526
XX ABS526
XX IS-NOV
DT 15-NOV
DT 15-NOV
DT Human
XX Human
XX
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Best Local !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a polynucleotide derived from the 4q12-4q13 region of human chromosome 4 and encoding a zsig63 polypeptide, a secreted salivary protein with anti-microbial activity. Due to their microbial activity, the sequences can be used in the study of microbial infections, e.g. for recombinant production of anti-microbial proteins. The sequences can be used in the treatment of tooth decay, periodontal disease, thrush, gastrointestinal disease, urinary tract infections, vaginal infections, skin infections, epithelial wounds, chronic tissue damage, acquired immunodeficiency syndrome (AIDS), cystic fibrosis, lung infections, sarcoidosis, emphysema and chronic bronchitis. This sequence represents a sequencing primer for cDNA encoding human zsig63
                                                                    Human; secreted salivary protein; zsig63; immunogen; zsig63-cytokine; antibody-cytokine, in vivo killing; pathological microbe; bacteria; fungal; viral; infection; salivary gland; anti-microbial; dental caries; tooth decay; periodontal disease; thrush; gastrointestinal disease; urinary tract infection; vaginal infection; skin infection; microflora; epithelial wound; pathogenic colonisation; invasion; pro-inflammatory; chronic tissue damage; vascular system; diabetes; anti-inflammatory; incompetent immune system; AIDS; acquired immunodeficiency syndrome; chemotherapy; radiation treatment; lung infection; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; zsig63; chromosome 4q12-4q13; salivary protein; antimicrobial; ss; microbial infection; tooth decay; periodontal disease; thrush; emphysema; gattrointestinal disease; urinary tract infection; vaginal infection; skin infection; epithelial wound; chronic tissue damage; cystic fibrosis; acquired immunodeficiency syndrome; AIDS; lung infection; sarcoidosis; chronic bronchitis; gene therapy; protein therapy; primer; ZC7231.
                                   chemotherapy; radiation digestion; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABS52637 standard; DNA; 26 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 26 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Polynucleotides encoding salivary proteins useful as anti-microbial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-MAR-2000; 2000US-00527345.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Adler DA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ZYMO ) ZYMOGENETICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                     salivary protein zsig63 PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 18.4; DB 1;
Pred. No. 3.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 26;
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RESULT 164
AAD45054/c
ID AAD450
XX
AC AAD450
AC AAD450
XX
DT 27-DEC
XX
DE ZC7231
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                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a new secreted salivary protein, zsig63. CC The invention is useful for detecting in a test sample, the presence of CC and so useful as an immunogen for producing an antibody to zsig63 cC golypeptide. zsig63-cytokine fusion proteins or antibody-cytokine fusion CC protein are useful for enhancing in vivo killing of target tissues. CC Pharmaceutical composition comprising purified zsig63 polypeptide are useful for enhancing in vivo killing of target tissues. CC Pharmaceutical composition comprising purified zsig63 polypeptide are useful in the treatment of conditions associated with pathological CC microbes, including bacterial, fungal and viral infections. High CC expression of zsig63 in salivary gland suggests that anti-microbial CC polypeptides are useful for treatment of dental caries (tooth decay), CC periodontal disease, thrush and gastrointestinal disease. Other CC applications can be used in urinary tract infections, vaginal infections, prevention of infection in skin and other epithelial wounds. The CC against pathogenic colonisation and invasion. The invention is useful CC when pro-inflammatory activity is desired. Applications for such pro-inflammatory activity is desired. Applications for insure pro-inflaminatory desired with dispersion is colonists of the invent
                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                    Matches
                                               27-DEC-2002
                                                                                                                                   AAD45054 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 26 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel secreted salivary protein, zsig63 and polynucleotide encoding it useful for treating microbial infections, inflammatory conditions, dental caries and lung infections associated with cystic fibrosis.
  2C7231 primer used in the identification of human
                                                                                        AAD45054;
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17-MAR-2000; 2000US-00527345.
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(SHEP/)
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SHEPPARD P
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                                           (first entry)
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Pred. No. 3
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CC as zaig63 and nucleic acid molecules encoding such polypeptides. zaig63 cc an be used in detecting agonists and antagonists of its activity, and is also useful as a host defense polypeptide, immune modulating factor, cc antipathogenic polypeptide, cell-cell signalling molecule, growth factor, cc cytokine, or as secreted extracellular matrix associated proteins with cc cytokine, or as secreted extracellular matrix associated proteins with cc growth factor hormone activity. It is useful for treating conditions consisted with pathological microbes, including bacterial, fungal and controlled through and gastrointestinal disease, for treating conditions context of infection, vaginal infection and for preventing infection in skin and context of lung dysfunction, salivary pland dysfunction and invasion, for treating chronic tissue damage e.g. damage in extremities associated with disease and useful as anti-inflammatory agents. It is useful as a marker of lung dysfunction, salivary gland dysfunction, or dysfunction of constate gland. It is also therapeutically useful for aiding digestion. Cc polynucleotides of the invention are used in gene therapy for increasing chromosome 4 associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as chromosome 4 associated with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease or other human traits and as consisted with disease disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel secreted salivary polypeptide, zsig63, useful as antimicrobial agent for treating microbial infection, dental carries, periodontal disease, thrush gastrointestinal disease, and for aiding digestion.
                                                                                                                                                                                             Sequence 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to human secreted salivary polypeptide designated as zsig63 and nucleic acid molecules encoding such polypeptides. zsig63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 29; 33pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ADLE/) ADLER D A.
(SHEP/) SHEPPARD P O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                forensic DNA profiling; chondrosarcoma; atherosclerosis; primer;
                                              5392 TAAAAAATACAAAAAAGAAAAAA 5415
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26
                                                                                                                       Similarity
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                                                                                                Conservative
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RESULT

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ABX93599/c
ID ABX935
XX ABX935
XX ABX935
XX ABX935
XX ABX935
XX ABX935
XX BBS; PC
KW BB; PC
KW BB; PC
KW Urinar
KW Chroni
KW Chroni
KW Chlon
KW Chl
CC least 90% identity to an amino acid sequence which comprises domain 1 of crisig63, domain 2, domain 3, mature zsig63 and full length zsig3 Also CC included are the polynucleotide encoding zsig63 and full length zsig3 Also CC included are the polynucleotide encoding zsig63 at zsig63 expression CC vector, a cultured cell comprising the vector and expressing the protein, and DNA encoding a fusion protein (comprising amino acids 1-15, 16-37, 38-CC 126, 127-219 or 16-219 of zsig63 and an additional protein), using a casig63 reporter gene construct to identify zsig63 agonists, and producing CC an anti-zsig63 antibody using zsig63 immunogenic peptides, Zsig63 is CC useful for detecting in a test sample, the presence of antagonist of CC zsig63 protein activity. Zsig63 sequence of antagonist of CC exhibits high expression in salivary gland, can be used for treating CC dental carries, periodontal disease, thrush, and gastrointestinal CC disease, urinary tract infections, vaginal infections, skin infections and other epithelial wounds. The polypeptides can be used to establish CC normal microflora and protect against pathogenic colonization and CC invasion. Zsig63 can also be used for providing pro-inflammatory activity for treating chronic, tissue damage particularly in areas having limited cordinates and protect against pathogenic colonization and CC immunocompromised AIDS patients or in individuals that have undergone constitered providers and protections e.g. in diabetus, and for treating lung discussions e.g. in diabetus, and for treating e.g. in CC levels in the trachea may indicate that such polypeptides may serve as a cC conditions associated with salivary gland or lung dysfunction including conditions bronic bronichtis, prostate dysfunctions such as prostate
CC adenocarcinoma, aiding digestion, and as components of defined cell culture media and may be used to replace serum that is commonly used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; PCR; zsig63; adhesin; salivary gland; dental carries; periodontal disease; thrush; gastrointestinal disease; epithelial wound; urinary tract infection; vaginal infection; skin infection; primer; pro-inflammatory; chronic tissue damage; vascular system; diabetes; AIDS; lung infection; cystic fibrosis; lung dysfunction; digestive; salivary gland carcinoma; Pneumocystis carinii infection; emphysema; chronic bronchitis; prostate dysfunction; prostate adenocarcinoma; cell culture media; gene therapy; human chromosome 4q12-4q13; dentinogenesis imperfecta; dentin dysplasia type II.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated zsig63 polypeptide, member of the adhesin family, useful for treating dental carries, periodontal disease, thrush, gastrointestinal disease, urinary tract infections, vaginal infections,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to an isolated zsig63 polypeptide comprising least 90% identity to an amino acid sequence which comprises domain
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(SHEP/) SHEPPARD P O.
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Query Match Best Local S Matches 20

Similarity

0.3%;

Conservative

1; Mismatches

Score 18.4; Pred. No. 3

DB 1;

Length 26

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Gaps

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ID ACF363
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                                                    electrophoresis, and as internal controls allowing for calculation of relative amounts of material present. The present sequence represents specific example of a PCR primer used in the method of the invention
                                                                                                                                                 The invention relates to producing a population of double-stranded product DNA molecules comprising amplification by a nested PCR method. The method is useful in profiling mENA transcribed in a system under investigation. The oligonucleotides are used as size standards in
   Sequence 26
                                                                                                                                                                                                                                                                                                   Claim 6; Page 85; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                   Producing a population of double-stranded product DNA for mRNA profiling, comprises amplification by nested
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Linnarsson S,
Montelius A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-618365/58
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    Length 26;

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RESULT 167
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Best Local
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                   G protein-coupled receptor; GPCR; drug screening; diagnosis; transgenic mouse; neurological disorder; adrenal gland disorder; colon disorder; intestinal disorder; cardiovascular disorder; muscular disorder; blood disorder; immune disorder; bone disorder; joint disorder; metabolic disorder; mutritive disorder; cancer; kidney disorder; liver disorder; lung disorder; breast disorder;
                                                                                                                                                                                                                                                                                                                         ADO30495;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a DNA molecule containing a sequence complementary to a Bovine Viral Diarrhea Virus (BVDV) RNA. The RNA when introduced into susceptible host cells, induces the generation of infectious BVDV particles. The attenuated BVDV clone or strain is useful in the manufacture of a vaccine for the prophylaxis and treatment of BVD infections. The invention is useful in gene therapy. The present sequencies a PCR primer used to amplify BVDV gene. (Updated on 27-OCT-2003 to
                                                                                                                                                                                                                                                                                                                                                                          ADO30495 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 26 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and treatment of Bovine Viral Diarrhea Virus (BVDV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-SEP-2001; 2001DE-01043813
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-SEP-2002; 2002WO-EP009925
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bovine viral
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA molecule useful for manufacturing a vaccine for the prophylaxis treatment of Bovine Viral Diarrhea Virus (BVDV) infections, comprise
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5392 TAAAAAATACAAAAAGAAAAAA 5415
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  disorder; uterus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                              PCR primer, SEQ ID NO:1598.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BAAAAAAAAAAAAAAAAAAAA 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Mismatches
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Pred. No. 3
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skin disorder; stomach disorder; pancreas disorder; spleen disorder; thymus disorder; thyroid disorder; antiparkinsonian; antimanic; cytostatic; antiinflammatory; vasotropic; antianginal; antiarrhythmic; CNS; central nervous system; respiratory; antidiarrhoeic; antidiabetic; virucide; hepatorropic; antibacterial; antidiarrhoeic; antidiabetic; dermatological; antiulcer; antithyroid; antiallergic; anorectic; immunosuppressive; nephrotropic; gene therapy; GPCR modulator; rapid amplification of cDNA ends; RACE PCR; primer; ss.

Synthetic.

WO2004040000-A2.

13-MAY-2004.

09-SEP-2003; 2003WO-US028226.

09-SEP-2003; 2003WS-0409303P.

09-APR-2003; 2003US-0461329P.

(PRIM-) PRIMAL INC.

Gaitanaris GA, Bergmann JE, Gragerov A, Hohmann J, Li F; Madisen L, Mcilwain KL, Pavlova MN, Vassilatis D, Zeng H;

WPI; 2004-390329/36.
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Novel mammalian G protein coupled receptors, useful for identifying compounds that modulates diagnosing and treating disease condition associated with GPCR dysfunction e.g. autoimmune diseases, angina pectoris, Parkinson's disease.

Disclosure; SEQ ID NO 1598; 542pp; English.

CC (GPCRs) and nucleic acids encoding them. The invention also relates to CC sequences at least 90% identical to the GPCR proteins and nucleic acids co of the invention; methods of treating, preventing or diagnosing diseases CC associated with GPCRs of the invention; methods of screening for CC compounds useful in the treatment of GPCR related diseases; a transgenic CC mouse comprising a GPCR gene of the invention; methods of screening for CC compounds useful in the treatment of GPCR-related diseases; a transgenic CC must comprising a GPCR gene of the invention; a mouse comprising a GPCR gene of the invention; and kits comprising cCC mouse comprising could discove the transgenic mice; kits comprising several mice, each of which has CCC probes which hybridise to GPCR polynucleotides of the invention. The CCC invention further discloses variants of the invention; and kits comprising CCC probes which hybridise to GPCR polynucleotides of the invention. The CCC comprising a GPCR nucleic acid. The GPCR nucleic acids and proteins may CCC used in the diagnosis, treatment or prevention of a wide variety of CCC diseases including neurological disorders (e.g., Alzheimer's disease, CCC diseases including neurological disorders (e.g., Alzheimer's disease, CCC diseases including neurological disorders of the colon or intestine CCC (e.g., Crohn's diseascular disorders (e.g., angina, cardiac arrhythmia or CCC myocardial infarction); muscular disorders (e.g., angina, cardiac arrhythmia or CCC myocardial pland; disorders (e.g., angina, cardiac arrhythmia or CCC myocardial pland; disorders (e.g., angina, cardiac arrhythmia or CCC myocardial pland; disorders (e.g., angina, cardiac arrhythmia or CCC myocardial pland; disorders (e.g., angina, cardiac arrhythmia or CCC myocardial pland; disorders (e.g., angina, cardiac arrhythmia or CCC myocardial pland; disorders (e.g., angina, cardiac arrhythmia or CCC myocardial pland; disorders (e.g., angina); disorders (e.g., angina); disorders (e.g., angina); disorders (e.g., angina); disorders

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Sequence 26 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 2 Other;
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Query Match 0.3%; Score 18.4; DB 1; Length 26; Best Local Similarity 83.3%; Pred. No. 3.4e+02; Matches 20; Conservative 1; Mismatches 3; Indels 0; Gaps 0;

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RESULT 170
AAC62451/c
ID AAC624
XX
AC AAC624
XX
DT 07-FEH
XX
DE Cleava
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AAC62450/c
ID AAC624
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XX AAC624
XX O7-FEB
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XX O5-OCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention is concerned with the cleavage of nucleic acids from solid supports. This is carried out by adding a non-conventional nucleotide into the nucleic acid attached to the support, so that it is recognised and cleaved by a specific DNA glycosylase and the sequence is released. This is useful in many molecular biological procedures such as sequencing, in vitro amplifications, CDNA and template preparation, DNA-based assays, mutagenesis procedures, nucleic acid purification and affinity chromatography. The present sequence is an oligonucleotide used in assays to demonstrate the methods of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 23 BP; 0 A; 0 C; 0 G; 22 T; 1 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detaching nucleic acid molecule comprising unconventional nucleotide incorporated at predetermined site from a solid support involves cleaving the nucleic acid molecule at the site of unconventional nucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-664908/64.
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   Cleavage of nucleic acids from
                                                                07-FEB-2001
                                                                                                                          AAC62451;
                                                                                                                                                                                    AAC62451 standard;
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87.0%;
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Pred. No. 3.6e+02;
      solid supports assay oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
         #2
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RESULT 171
AAT99286
ID AAT992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local S
Matches 20
                                     28-AUG-1992;
02-FEB-1993;
06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention is concerned with the cleavage of nucleic acids from solid supports. This is carried out by adding a non-conventional nucleotide into the nucleic acid attached to the support, so that it is recognised and cleaved by a specific DNA glycosylase and the sequence is released. This is useful in many molecular biological procedures such as sequencing, in vitro amplifications, CDNA and template preparation, DNA-based assays, mutagenesis procedures, nucleic acid purification and affinity chromatography. The present sequence is an oligonucleotide used in assays to demonstrate the methods of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid cleavage; solid support; affinity chromatography; sequencing; mutagenesis; DNA preparation; nucleic acid purification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detaching nucleic acid molecule comprising unconventional nucleotide incorporated at predetermined site from a solid support involves cleaving the nucleic acid molecule at the site of unconventional nucleotide.
                                                                                            07-JUN-1995;
                                                                                                                       30-SEP-1997
                                                                                                                                                 US5672479-A.
                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                   cancer; probe;
                                                                                                                                                                                                                                  PUR element; human;
                                                                                                                                                                                                                                                                                       15-APR-1998
                                                                                                                                                                                                                                                                                                                                            AAT99286 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 23 BP; 0 A; 0 C; 0 G; 0 T; 23 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 32; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-664908/64.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-MAR-2000; 2000WO-GB001190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                           Homo sapiens.
                                                                                                                                                                                                                                                            POLYA, a competitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GOLD/) GOLDSBOROUGH
                                                                                                                                                                                                                                                                                                                                                                                                                                         5393 AAAAAATACAAAAAGAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                               23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                               AAAAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                       (first
                                      92US-00938189.
93US-00014943.
95US-00470911.
                                                                                                                                                                                                                   hybridisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99GB-00007245
                                                                                            95US-00486421.
                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                 c-myc; inhibitor; hyperproliferative
                                                                                                                                                                                                                                                            oligonucleotide for binding human PUR-alpha.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.3%;
                                                                                                                                                                                                                                                                                                                                             BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 18.2; DB 1;
Pred. No. 3.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 23;
                                                                                                                                                                                                                                    disease; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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0,

(MOUN) MOUNT

SINAI SCHOOL MEDICINE

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RESULT 172
AAV31743
ID AAV317
AC AAV317
XX AV317
XX Nucleo
XX PUR-al
KW PUR-al
KW Hyperp
XX Synthe
PN US5756
XX US5756
XX O6-JUN
XX PR 02-FEB
XX PR 02-FEB
XX PR 02-FEB
XX PR PT PUR PX
PT
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     Query Match
Best Local S
Matches 20
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Best Local S
Matches 20
                                                                                                                                                               This is the nucleotide sequence of an oligonucleotide used as a competitor with the PUR element in the method of the invention, involving the use of the PUR protein and its fragments, which inhibit PUR protein binding to PUR element or other proteins. Inhibitors of PUR activity may be useful for treating viral infections and hyperproliferative diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The oligonucleotides AAT99279-T99286 were used as competitor oligonucleotides for the binding of PUR preein to DNA. The PUR sequence can be used to identify chemical or biological compounds that bind to PU or binding fragments of PUR. Inhibitors of PUR activity may be used to treat hyperproliferative diseases such as cancer
                                                                                                Sequence 24 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Assays for protein or
                                                                                                                                                                                                                                                                                                                    Example 7.1.1; Col 33; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                              PUR protein and its fragments - that inhibit PUR protein binding to
                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-321632/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bergemann AD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-AUG-1992;
02-FEB-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US5756684-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PUR-alpha gene; inhibition; hyperproliferative disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleotide sequence of the oligonucleotide POLYA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV31743;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV31743 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (MOUN ) MOUNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5393 AAAAAATACAAAAAGAAAAA 5415
     l Similarity
20; Conserv
                                                                                                                                                  cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                     or other proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Col 33; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAAAAAAAAAAAAAAAAAAA 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ð
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fragments, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SINAI SCHOOL MEDICINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            92US-00938189
93US-00014943
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-00470911.
                                                                                                  24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
                         0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ligands or modulators - using immobilised PUR cotreat hyper-proliferative diseases, e.g. can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <u>묒</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24
; Score 18.2; D; Pred. No. 3.6e 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              viral infection; cancer;
ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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Pred. No. 3.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                           .6e+02
                                                  DB 1; Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
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       Indels
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     0
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     Gaps
                                                                                                                                                                                                                                                involving
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RESULT 174
AAA40359/c
ID AAA403
XX
AC AAA403
XX
AC AAA403
XX
TO 10-NOV
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DE pBlues
XX
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Best Local S
Matches 20
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02-FEB-1993;
06-JUN-1995;
                                                                                                                                                                                                                                                          The present invention describes a monoclonal antibody that specifically binds to an epicope of the PUR protein. Antibodies that bind to the PUR protein and neutralise PUR activity may be used to treat hyperproliferative diseases such as cancer. PUR antibodies may be used diagnostically to detect aberrant expression of the PUR protein and/or mutations in the PUR gene. The present sequence represents an oligonuclectide used in the cloning and sequencing of the PUR protein and its sequence element PUR repeat, in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAX04086 standard; DNA;
            pBluescriptSK+
                                   10-NOV-2000
                                                                                                                                                                                                                                        Sequence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide POLYA used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX04086;
                                                           AAA40359
                                                                                 AAA40359 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-152881/13
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bergemann AD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PUR element; PUR-alpha; hyperproliferative disease; cancer; monoclonal antibody; identification; characterisation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-APR-1999
                                                                                                                                                                                                                                                                                                                                                                    Example; Col 33; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                         Monoclonal antibody specific for PUR protein - useful for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MOUN ) MOUNT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           09-FEB-1999
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                                                                                                                                                                                                      Similarity
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                                                                                                                                           AAAAAAAAAAAAAAAAAAAA 23
                                                                                                                                                                                          Conservative
                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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93US-00014943.
95US-00470911.
           phagemid
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Johnson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-00486809.
                                                                                  RNA;
                                                                                                                                                                                                    0.3%;
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            primer
                                                                                  24
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                                                                                                                                                                                                      Score 18.2;
Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PUR cloning and sequencing
             SEQ ID
                                                                                                                                                                                           Mismatches
              NO:
                                                                                                                                                                                                      .6e+02
                                                                                                                                                                                                                 DB 1;
              9
                                                                                                                                                                                                                Length 24;
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ARAGONIT 175
ARAGON 53/C
ID ARAGON
XX ARAGON
AC ARAGON
AC ARAGON
AC ARAGON
AC PBlues
XX PBlues
XX PBlues
XX Primer
XX Primer
XX WO2000
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XX IT-DEC
XX IT-DEC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes a novel method (M1) for inserting a nucleic acid (N1) into a circular vector (V1) comprising joining ends of N1 and V1 under a first nucleic acid concentration, melting hybridized cohesive circularization ends, and reannealing the ends at a second concentration. The methods are useful for the cloning small amounts of nucleic acids and forming genomic libraries of complex populations of pNA or cDNA. The methods allow the cloning of minute amounts of nucleic acids efficiently and avoids the size selection problems of prior art systems. Larger nucleic acid fragments are just as easily cloned, allowing highly representative libraries to be made. Vector to vector ligation is avoided using the methods. AAA40365 represents primers used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Inserting a nucleic acid into a circular vector comprising joining their ends, melting, and reannealing ends at two different concentrations, useful for cloning small amounts of nucleic acids and forming genomic
                                                                                                                                                                                                                                  pBluescriptSK+ phagemid primer SEQ ID
                                                                                                                                                                                                                                                                 10-NOV-2000
                                                                                                                                                                                                                                                                                                                             AAA40353 standard; DNA; 24 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24 BP; 0 A; 0 C; 0 G; 16 T; 8 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2000-442381/38
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                                              17-DEC-1998;
                                                                                                           22-JUN-2000.
                                                                                                                                          WO200036088-A1
                                                                                                                                                                                                    Primer; cloning; ligation; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Romantchikov Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200036088-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Primer; cloning; ligation; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ROMA/) ROMANTCHIKOV Y.
                 (ROMA/) ROMANTCHIKOV Y
                                                                                                                                                                                                                                                                                                                                                                                                                        5393 AAAAAAATACAAAAAAAGAAAAAA 5415
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Similarity 87.0%;
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                                              98US-00213834.
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                                                                             99WO-US030277.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 18.2; DB 1;
Pred. No. 3.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Indels
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RESULT 176
AAP99756/c
ID AAP9977
XX AAP9977
XX AAP997
XX Vaccin
KW Vaccin
KW immuno
KW infect
XX Synthe
XX W02001
XX W02001
XX E-SEP
PR 25-SEP
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PA
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27-SEP-1999;
23-AUG-2000; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                under a first nucleic acid concentration, melting hybridized cohesive circularization ends, and reannealing the ends at a second concentration. The methods are useful for the cloning small amounts of nucleic acids and forming genomic libraries of complex populations of DNA or CDNA. The methods allow the cloning of minute amounts of nucleic acids efficiently and avoids the size selection problems of prior art systems. Larger nucleic acid fragments are just as easily cloned, allowing highly representative libraries to be made. Vector to vector ligation is avoided using the methods. AAA40351-A40366 represents primers used to illustrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-APR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Immunostimulatory nucleic acid #872.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF99756 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes a novel method (M1) for inserting a nucleic acid (N1) into a circular vector (V1) comprising joining ends of N1 and V1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 66;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Romantchikov Y;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAAAAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                      Schetter C,
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99US-0156135P
2000US-0227436P
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                                                                                                                                                                                                                         Vollmer
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Pred. No. 3.6e+02
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Claim 101; Page 57; 338pp; English

Vaccinating against tumors, infectious diseases, allergies using immunostimulatory Py-rich and TG nucleic acids.

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asthma

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RESULT 177
AAF99304/c
ID AAF993
XX AAF993
XX Immuno
XX Immuno
XX Vaccin
KW infect
XX Vaccin
KW infect
XX W02001
XX W02001
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Best Local S
Matches 20
immune response. The present sequence acid. The immunostimulate (py-rich) or thymidine (T) rich. against tumour antigene ......
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                                                                                                response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory
                                                                                                                                                                                                                                                                                                                                                 Vaccinating against tumors, infe using immunostimulatory Py-rich
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-SEP-199; 99US-0156113P.
27-SEP-199; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
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ponse. The method comprises administering an immunostimulatory
d to a non-rodent subject in sufficient quantity to stimulate a
une response. The present sequence is one such immunostimulator
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                                                                                                                                                                                                                                                                             Page
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schetter C,
                                                                                                                                                                                                            invention relates to a method for stimulating
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   immunostimulatory nucleic acids can be pyrimidinė rich
idine (T) rich. The method is used to vaccinate subjects
tigens, viral antigens (e.g. herpesviridae, retroviridae
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                                                                                                                                                                                                                                                                                                                                                     infectious diseases, allergies and asthmarich and TG nucleic acids.
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RESULT 178
AAF99757
ID AAF99757
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XX AAF9977
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Matches 20
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27-SEP-1999;
23-AUG-2000; 20
                                                         response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parassitic antigens. The method is staphylococcus), fungal antigens and/or parassitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vaccinating against tumors, infectious diseases, allergies and using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 101; Page 57; 338pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-JUN-2001
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                            present sequence may have a phosphorothioate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-pimmunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                      present invention relates to a method for stimulating ponse. The method comprises administering an immunosting
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Schetter C,
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99US-0156135P.
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Pred. No. 3
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Matches 20
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Matches 20; Conserv
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16-MAR-2000; 2000US-0189862P.
25-MAY-2000; 2000US-0207454P.
09-JUN-2000; 2000US-0211314P.
18-JUL-2000; 2000US-0219007P.
13-DEC-2000; 2000US-0255281P.
                                                                                                                                                                                                                                                               The invention relates to an isolated nucleic acid molecule (I) comprising a nucleotide sequence given in Tables 1-9 (ABV00010-ABV62213) of the specification or its complement. (I) is useful for: (a) assessing whether a patient is afflicted with prostate cancer; (b) monitoring the progression of prostate cancer in a patient; (c) assessing the efficacy of a test compound to inhibit prostate cancer in a patient; (d) assessing the efficacy of a thorapy for inhibiting prostate cancer in a patient; (e) selecting a composition for inhibiting prostate cancer in a patient; (f) assessing the prostate cell carcinogenic potential of a compound; (g) determining whether prostate cell carcinogenic potential of a compound; (g) determining whether prostate cancer has metastasized in a patient; (h) assessing the aggressiveness or indolence of prostate cancer in a patient; (l) is also useful as a pharmacodyanamic or pharmacogenomic marker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel isolated nucleic acid molecule associated with cancerous state prostate cells and correlating with presence of prostate cancer, usefor detecting presence of prostate cancer, stage of prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; prostate cancer; cytostatic; carcinogen; pharmacodyanamic marker; pharmacogenomic marker; gene; ss.
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                                                   5393 АЛАЛАЛАТАСАЛАЛАДАЛАЛАЛ 5415
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                                                                                                          0, Mismatches
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Pred. No. 3.6e+02
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Pred. No. 3.6e+02;
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                                                                                                                                                                                                                                                                                                            CC The invention relates to inhibiting angiogenesis in a subject, comprising CC administering at least one antiangiogenic nucleic acid molecule. Also CC included is a kit comprising a first container housing the antiangiogenic CC nucleic acids, and instructions for administering them to a subject chaving a condition characterised by unwanted angiogenesis. The method is CC useful for inhibiting angiogenesis associated with solid tumour growth, CC tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, CC diabetic retinopathy, retinopathy, retropathy, retrolental fibroplasia, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, CC rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque Cc neovascularisation, intestinal adhesions, atheroseis, scleroderma and Cc hypertrophic scars. The present sequence is an antiangiogenic nucleic acid of the invention
                                                                                                                         Query Match
Best Local S
Matches 20
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                                                                                                                                                                                                                                                      Sequence 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Angiogenesis inhibitory oligonucleotide #961
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                                                            5393 AAAAAATACAAAAAGAAAAAA 5415
24
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                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                      BP; 0 A; 0 C; 0 G; 24 T; 0
                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
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                                                                                                                         0
                                                                                                                      Score 18.2; DB 1
Pred. No. 3.6e+02
0; Mismatches
                                                                                                                                                                                                                                                             U; 0 Other;
                                                                                                                                                                                         DB 1; Length
                                                                                                                             Indels
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                                                                                                                             Gaps
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RBSULT 181
ABS77949/c
ID ABS77949 standard; D
XX
AC ABS77949;

DNA;

24

BP.

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RESULT 182
ABS78478
ID ABS784
XX
AC ABS784
XX
DT 13-DEC
XX
DE Angiog
XX
ANGIOG
KW Angiog
KW Lumour
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Best Local
                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasts, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic acid of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiettasia; haemophiliac joint; plaque neovascularisation; telangiettasia; haemophiliac joint;
Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis;
                                                                                                                                                                                                                                                                                                                                                            Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Inhibiting angiogenesis in a subject, involves admin antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2002-566690/60.
                                               Angiogenesis inhibitory oligonucleotide #962.
                                                                                13-DEC-2002
                                                                                                                                                ABS78478 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    14-DEC-2001; 2001WO-US048458
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-DEC-2002
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                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                AAAAAAATACAAAAAAAGAAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Page 27; 276pp; English
                                                                                                                                                                                                                                 AAAAAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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Pred. No. 3
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 RESULT 183
ABL39405/c
ID 39405/c
ID 394294
XX ABL394
XX ABL394
XX Inmuno
XX Inmuno
XX Antibo
XW Antibo
XW angiog
XX Synthe
XX Synthe
XX EH Key
FT modifi
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16-APR-2002

(first entry

ABL39405;

Immunostimulatory nucleic acid

SEQ ID NO: 841.

Antibody-induced cell lysis; cancer; immunostimulatory; angiogenesis; metastasis; cytostatic; phosphorothioate

; CD20; backbone;

88

modified_base

1. .24 /*tag= a /mod_base= OTHER Location/Qualifiers ABL39405 standard; DNA; 24 BP

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Matches
                                                                                                                            Query Match
Best Local (
                                                                                                                                                                                                                                                                               administering at least one antianglogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antianglogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted anglogenesis. The method is useful for inhibiting anglogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myccardial anglogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, anglofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diabetic retinopathy; retinopathy of prematurity; macular degeneration corneal graft rejection; neovascular glaucoma; retrolental fibroplasia rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Inhibiting angiogenesis in a subject, involves administering antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    scleroderma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2002-566690/60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention
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                                                                                                                                  Similarity
                                                                                                                                                                                                             24
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AAAAAAAAAAAAAAAAAA 23
                                                                                                      Conservative
                                                                                                                                                                                                             BP; 24 A; 0 C;
                                                                                                                                                                                                                                                                    invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypertrophic scar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  36; 276pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 relates to inhibiting angiogenesis in
                                                                                                                            0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  English.
                                                                                                                                                                                                             0
                                                                                                   Pred. No. 3.60
0; Mismatches
                                                                                                                               Score 18.2; DB 1;
Pred. No. 3.6e+02;
                                                                                                                                                                                                             G; 0 T; 0 U;
                                                                                                                                                                                                               0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ty; macular degeneration; retrolental fibroplasia;
                                                                                                                                                        Length 24;
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                                                                                                         Gaps
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RESULT 184
ABX14631/C
ID ABX146
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ABX14631/C
AC ABX146
XX
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AX
OGAMAR

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Best Local Sim.
Matches 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for the cell surface antigens and antibodies are such as basal cell carcinoma, bladder treating or preventing cancer such as basal cell carcinoma, bladder treating or preventing cancer such as basal cell carcinoma.
                                                                                                                                                                                                                                                                                                                              es; guanosine triphosphatase activator protein 10.12; PCR; primer; malignant tumour; inflammation; immunological disease; haemopathy; human immunodeficiency virus infection; HIV; RT-PCR; reverse transcriptase PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Guanosine triphosphatase activator protein 10.12 RT-PCR primer #1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABX14631;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 309; 312pp; English
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      10-NOV-2000, 2000CN-00127330
                                                                    10-NOV-2000; 2000CN-00127330.
                                                                                                                                                                                                         CN1352022-A.
                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABX14631 standard; DNA; 24 BP
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Pred. No. 3.
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RESULT 185
ABA98840
ID ABA988
XX ABA988
XX ABA988
XX OTHER
DT 01-JUL
XX Compon
KW metabo
XX Synthe
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YT modifi
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Best Local S
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Detection of multiple analytes, and pollutants, involves adding
                                                WPI; 2002-164078/21.
                                                                                                                                             04-MAY-2000; 2000US-00564230
                                                                                                                                                                            03-MAY-2001; 2001WO-US014528
                                                                                                                                                                                                            08-NOV-2001
                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                    Component detection; clinical diagnosis; cell detection; drug detection; metabolite detection; pesticide detection; ligand detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  A24 oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 24 BP; 5 A; 5 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New polypeptide-quanosine triphosphatase activator protein 10.12 polynucleotide for encoding such polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-714410/78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                мао Y,
                                                                                                                                                                                                                                           WO200184157-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BODE-) BODE GENE DEV CO LTD SHANGHAI.
                                                                                                                (DADE-) DADE BEHRING INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3594 TGCTCAGGCTAATCTCAAACTCC 3616
                                                                               JS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24 TGCCCAGGCTGGTCTCAAACTCC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Page 17
                                                                              Cromer R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
ilarity 87.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                  /*tag= a
/label= OTHER
/note= "modified by PO2OCH2CH2CH2SSCH2CH2CH2OH"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (disclosure); 33pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                      for the creation of Pc-A24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                 Patel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ₽P
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                                                                              ₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 18.2;
Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
   e.g. ligands, receptors, polynucleotides a combination of sensitizer reagents and
                                                                                 Kurn
                                                                               z
                                                                                 þ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
                                                                                 Keczer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                 S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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RESULT 186
AAS17869
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                              Detecting presence of polynucleotide, differences between polynucleotide sequences, useful for detecting single nucleotide polymorphism and alleles of polynucleotide sequence involves use of three competitive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphism detection; sequence detection; mutation detection; A24; probe; non-dissociative termolecular complex; dopTAR sensitiser particle; single nucleotide polymorphism; SNP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAS17869 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                probes.
                                                                                                    WPI; 2002-097664/13.
                                                                                                                                                                                                        19-MAY-2000; 2000US-00574596
                                                                                                                                                                                                                                          17-MAY-2001; 2001WO-US016089
                                                                                                                                                                                                                                                                                                              WO200190399-A2
                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A24 oligonucleotide used to create dopTAR chemiluminescer particles.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         reactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5393 AAAAAATACAAAAAGAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   h 0.3%;
Similarity 87.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       reagent Actuatable by a product of the sensitizer reagents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAAAAAAAAAAAAAAAAA 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                       BEHRING INC
                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers 24
                                                                                                                                                                                                                                                                                                                                      /*tag= a
/note= "A is covalently linked to a
/note= "A is covalently linked to a
P020CH2CH2CH2CH2SSCH2CH2CH2OH moiety"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 18.2; DB 1;
Pred. No. 3.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3.6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 24;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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PX PPPX PX PX PX PX PX PX PX SX XX XX EXX PX AX X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention represents a method for detecting the presence of a CC polymucleotide sequence, differences in polymucleotide sequences or CC mutations in genomic DNA. The method involves contacting 3 contacting 3 polymucleotide probes with a sample containing a polymucleotide. The CC clist probe hybridises to a region of the polymucleotide sequence and the CC sequence. The second and third probes are identical except for the CC presence or difference of one or more nucleotides. The reaction medium is then subjected to conditions for forming substantially non-dissociative termolecular complexes, which can be at least one of, the polymucleotide sequence with the first and second probes or the polymucleotide sequence CC with the first and third probes. The oligonucleotide probes have labels con-covalently bound to allow for their detection upon binding. The CC mucleotide polymorphism (SNP) in a fragment of genomic DNA. The method cc and be used for the direct detection of mucleic acid in very small CC quantities without amplification. In addition, the method may be carried cout with amplification of the target and reference sequences. This sequence represents an oligonucleotide probe A24 used to create dopTAR CC binding the nucleic acid to a suspendable particle acts as a support and court of such a means of segregating the bound polynucleotide target from the court of the cou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local (
                                      Generating mRNA-cDNA hybrids for suppressing cancer-related genes, or treating or preventing microbe related genes, comprises thermocycling steps of promoter-linked double-stranded cDNA or RNA synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RNA-PCR; primer; ss; poly(dT)24; cytostatic; antibacterial; gene therapy; mRNA-DNA hybrid; gene function inhibition; cancer; PTGS; antisense; high throughput screening; D-RNA; DNA-RNA interference; RRAP; RNA dependent RNA polymerase; posttranscriptional gene silencing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 24 BP; 24 A;
                                                                                                                                       WPI; 2002-188740/24.
                                                                                                                                                                                                                                                                                                                          02-AUG-2001; 2001WO-US024412.
                                                                                                                                                                                                                                                                                                                                                                        07-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA-PCR procedure primer poly(dT)24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABK15639;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABK15639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example; Page 47; 75pp; English
                                                                                                                                                                                                                                                                            02-AUG-2000; 2000US-0222479P.
                                                                                                                                                                                                                                                                                                                                                                                                                    WO200210374-A2
                                                                                                                                                                                                                                 (UYSC-) UNIV SOUTHERN CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5393 AAAAAAATACAAAAAAGAAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            solution
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                Chuong C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry
                                                                                                                                                                                     Widelitz
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 18.2; DB 1,
Pred. No. 3.6e+02;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              G; 0 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Example 5; Page 26; 53pp; English

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RESULT 188
ADG16129/c
ID ADG161
XX ADG161
XX ADG161
XX Compou
XX Compou
XX Compou
XX Compou
XX Compou
XX W Phenot
XX W Phenot
XX Uniden
XX Uniden
XX Uniden
XX Uniden
XX W02001
XX W26-APF
PR 26-APF
PR 26-APF
PR 20-FEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to generating mRNA-cDNA hybrids, comprising (a) CC primers complementary to the sense conformation of the nucleic acid template, one or more contended one or more promoter-linked primers complementary to the grometer complementary to the contended one or more promoter-linked primers complementary to the contended contended contended one or more promoter-linked primers complementary to the promoter, (b) treating the nucleic acid template with an RNA CC primers to synthesise a first cDNA strand, (c) treating the first cDNA CC linked double-stranded nucleic acid, (d) treating the first cDNA CC linked double-stranded nucleic acid, (d) treating the promoter-linked CC complements. The method is useful for preparing high amounts of pure and CC specific mRNA-cDNA hybrids for transcription of the amplified mRNA fragments. The method is useful for preparing high amounts of pure and CC special contended in vivo, for inhibiting gene function in prokaryotes of vitro as well as in vivo, for inhibiting gene function in prokaryotes in vitro as well as in vivo, for inhibiting of gene function in prokaryotes of molecules, in high throughput screening of gene function in prokaryoted condidate molecular pathways with systematic knock out of involved conditions. The mRNA-cDNA hybrids may be used to screen for special gene functions, for manipulating gene expression in vitro, and constraints of the stability and effectiveness of transfected conscious activities. The RdRp (RNA dependent RNA polymerase) enzyme may consider a finity of the mRNA template of a D-RNA compared to dependent RNA polymerase) enzyme may consider the constraints of the sense-RNA template, resulting in multiple gene function. Sense than RNA-CRNA duplexes than RNA-CRNA duplexes. The cDNA part of a D-RNA compared to defect the sense of the sense-RNA template, resulting in multiple gene functions of the sense action optimizes to the present sequence of the transcribe mRNA interference effects with one probe. The present sequence is a polydly p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                compound activity characterisation; cellular activity; phenotypic attribute; candidate medicine; candidate treatment; multiple biological descriptor; cell marker; ss.
                                                                                                                                                                                                                                                                                                                                                      Compound activity characterisation-related oligonucleotide SeqID4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 24 BP; 0
                           26-APR-2000; 2000US-0199778P.
20-FEB-2001; 2001US-00790214.
                                                                                        24-APR-2001; 2001WO-US013248.
                                                                                                                                                                                                                         Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG16129 standard; DNA;
                                                                                                                                                                              WO200181895-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5393 АЛАЛАЛАТАСАЛАЛАДАЛАЛАЛ 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAAAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; 0 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 18.2; D. Pred. No. 3.6e. 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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S
                                                                                                                           Query Match
                                                                                                                                                            This invention relates to a novel method for the characterisation of the activity of a compound on cell. The method involves receiving images of cells with a cellular activity and images of other cells treated with the compound, quantitatively characterising phenotypic attributes of the image of cells with a cellular activity to produce a target phenotype for the cellular activity and that of the image of other cells to produce a second phenotype for the compound, and comparing the two phenotypes to determine whether the compound possesses cellular activity. The invention may be useful for characterising cellular activity of a compound, for determining information about properties of substances based upon the information about structure of living or non-living cells exposed to substances. The invention is also useful for identifying promising candidates in a search for new and better medicines and treatments using multiple biological descriptors from a single cell markers or components.
                                                                                                                         Sequence 24 BP; 0 A; 1 C; 0 G; 23 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Characterizing cellular activity of compound, by receiving images of cells with known activity and images of cells treated with compound, characterizing phenotypic attributes of images and comparing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Fig 18; 139pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oestreicher DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (CYTO-) CYTOKINETICS INC.
                                                            Local
    5393 AAAAAAATACAAAAAAGAAAAAA 5415
                                                                 Similarity
                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sabry JH,
                                                            0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Adams CL,
                                           0
                                         Pred. No. 3.60
0; Mismatches
                                                              Score 18.2;
Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Vaisberg EA,
                                                                 .6e+02;
                                                                                  DB 1; Length 24;
                                             ω.
                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Crompton
                                             <u>.</u>
                                             Gaps
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ABX79809/c
ID ABX798
XX
AC ABX798
AC ABX798
AC ABST; e
XX
DT 17-APR
XX
EST; e
KW EST; e
KW Predre
KW Fredre
KW GPINA1
AX
US6472
XX
US6472
XX
US6472
XX
PI 31-DEC
XX
PI 31-DEC
XX
PI 31-DEC
XX
PI GAZNE1
XX
PI GAZNE1
XX
PI GAZNE1
XX
PI Identi 밁 RESULT 189 EST; expressed sequence tag; ss; polymorphic repeat; tandem repeat; polymorphic marker prediction of ubiquitous simple sequences; POMPO Rep-K; human; genetic disease; nug-treatment; Machado-Joseph; Haw River syndrome; Huntington's disease; fragile-X syndrome; Fredreich's ataxis; myotonic dystrophy; hyperandrogenaemia; spinal atrophy; bulbar atrophy; spinocerebellar ataxia. ABX79809 US6472154-B1 Homo sapiens EST polymorphic DNA repeat polynucleotide #134. 17-APR-2003 ABX79809 standard; cDNA; 24 24 AAAAAAAAAAAAAAAAAA 2 (first entry) ВÞ

0;

Identifying a candidate polymorphic repeat within a coding sequence,

WPI; 2003-208818/20

Garner HR,

Wren JD,

Minna JD,

Fondon

31-DEC-1999; 31-DEC-1999; 29-OCT-2002

99US-00475947 99US-00475947.

(TEXA) UNIV TEXAS SYSTEM

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RESULT 190
ABZ8011
XX ABZ8011
AC ABZ801
XX IMMUNO
XX IMM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC The invention discloses a method for identifying a candidate polymorphic CC repeat within a coding sequence (expressed sequence tag, EST), which CC comprises detecting tandem repeats in a target coding sequence, scoring CC the repeats for polymorphic probability and generating a dataset CC correlating the repeats with polymorphic probability to identify a CC candidate polymorphic repeat. The computational methods (polymorphic CC marker prediction of ubiquitous simple sequences, POMPOUS, and Rep-X) are CC useful for identifying and detecting candidate polymorphic repeats in CC thuman genes, which can be used to understand, treat or eliminate genetic CC diseases, predispositions or adverse drug-treatment reactions. Examples CC of diseases linked to nucleotide repeats are Machado-Joseph, Haw River Syndrome, Huntington's disease, fragile-X syndrome, Fredreich's ataxis, CC myotonic dystrophy, hyperandrogenaemia, spinal and bulbar attrophy and CC spinocerebellar ataxia. The sequences presented in ABX79676-ABX80022 are CC the polymorphic repeats identified for a search of human ESTs
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Best Local S
Matches 20
                                                                 WPI; 2003-268241/26.
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03-JUL-2002; 2002US-0393952P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immunostimulation; immune response; natural killer cell; interferon; type 1 interferon; IFN; cancer; infectious disease; allergic disorde immune related disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABZ80181;
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                                                                                                                                                                             (COLE-) COLEY PHARM GROUP INC. (COLE-) COLEY PHARM GMBH. (IOWA ) UNIV IOWA RES FOUND.
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   immunostimulatory nucleic
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Pred. No. 3
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Example 1; Page 44; 115pp; English

for treating an allergic condition

The present invention describes immunostimulatory nucleic acids of 14-100 cc nucleotides in length comprising the formula 5' XIDCGHZ 3' (I), where XI cc x X2 = independently any sequence 0-10 nucleotides; D = nucleotide other CC immunostimulatory nucleic acid further comprises a sequence consisting of CC p and N positioned immediately 5' to X1 or 3' to X2 and N is a B cell CC neutralising sequence, where N begins with a CGG trinucleotide and is at CC least 10 nucleotides long and P is GC-rich palindrome containing sequence CC at least 10 nucleotides long. Also described: (1) a pharmaceutical CC composition comprising the immunostimulatory nucleic acid and a carrier; CC and (2) treating an allergic condition. (1) has antiallergic activity and CC can be used in gene therapy. (1) can be used for preparing a composition of cortrating a variety of immune related disorders such as cancer, CC infectious diseases and allergic disorders. (1) also stimulates the CC cactivation of natural killer cells and the production of type 1 cc oligonucleotide, which is used in an example from the present invention

Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 ₽; 0 Other;

Query Match Best Local S Matches 20 Similarity Conservative 0.3%; 0; Score 18.2; Pred. No. 3. Mismatches .6e+02; DB <u>,,</u> Length Indels 24; 0 Gaps 0,

RESULT 191
ACA622
ID ACA622
XX ACA622
XX ACA622
XX Oligo
XX Synthe
XX WENA e
XX US2003
XX Synthe
YN US2003
XX Synthe
YN US2003
XX Synthe
YN US2003
XX INDCOMPT Lin S,
XX INDCOMPT Lempli
PT acadia
PT scandia
PT Oligo (dT)24 RT-PCR primer. ACA62284; ACA62284 standard; 12-AUG-2003 (first entry) DNA; 24 ₽₽

PCR; PCR; RT-PCR; expression p ; primer; reverse transcriptase PCR; antisense therapy; profile; promoter containing primer.

Synthetic

US2003022318-A1

30-JAN-2003

07-SEP-2001; 2001US-00949305.

25-JAN-2000; 2000US-00494212

(EPIC-) EPICLONE

ß Ying S

WPI; 2003-479488/45

Improved polymerase thermocycling reaction for nucleic acid amplification, by thermal cycling of promoter-linked nucleic acid template synthesis and in vitro transcriptional amplification of promoter synthesis and in vitro transcriptional amplification of promoter synthesis. nucleic

Example 7; Page 14; 28pp; English

The invention relates to an improved polymerase thermocycling reaction (M1) for linear amplification of nucleic acid sequences, involves denaturing a number of nucleic acid templates (I), combining the denaturing a number of nucleic acid templates (I), combining the denatured (I) with a promoter-containing primer (P1), a primer (P2), a

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RESULT 192
ACD99729/c
ID ACD997
XX ACD997
XX ACD997
XX ACD997
XX Immuno
DE Immuno
XX Inmuno
XX Immuno
XX I
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determining protein expression of a cell in a diseased state, determining

the mRNA sequence for the aberrant proteins, synthesising an antisense
sequence of the mRNA, amplifying the antisense mRNA sequences by M1, and

delivering a pharmaceutically effective dosage of a composition

comprising the anti-sense mRNA and a compatible lipid based biological

corrier. M1 is also useful for predicting the efficacy of a proposed drug

targeted against an aberrant protein, by determining aberrant protein

production of cell in a diseased state by the above method, amplifying

the aberrant protein by M1 and using recombinant techniques to determine

the effect of proposed drug on the aberrant protein. M1 is also useful

for differential screening of tissue-specific gene expression at a

cellular level, for preparing labeled RNA/DNA probes for a gene chip

technology, and for determining the efficacy of a drug regiment against a

gene or its cDNAs. The present sequence is an Oligo (dT)24 RT-(reverse

transcriptage) PCR primer used to produce first strand cDNA in the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                         Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACD99729 standard; DNA; 24 BP
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                                                                                                                  Synthetic
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antilicer; gene therap; vaccine; non-allergic inflammatory disease psoriasis; ezzema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss
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                                                                                    WPI; 2003-521815/49.
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BERG D J.
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Pred. No. 3
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Treating non-allergic inflammatory diseases, allergic contact dermatitis, latex dermatitis

such as psoriasis, ecz
s or inflammatory bowel

immunostimulatory nucleic acid

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RRSULT 194
ACHO3284/c
ID ACHO322
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Best Local Similarity
Matches 20; Conserv
                                                          The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease.
                                                                                                                                                                                                                                                                                                                                                                                                             Treating non-allergic inflammatory diseases, such as psoriasis, ecz allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-521815/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-MAR-2001; 2001US-0279642P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACH03284;
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                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 34; 229pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (KRIE/) KRIEG A M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5393
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Pred. No. 3.6e
0; Mismatches
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Sequence 24

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comprises primer extension by reverse transcriptase and hybridising an coligonuclectide to the single stranded DNA, where the oligonuclectide is blocked from extension at the 3' end. The method is useful for amplification of nucleic acids. In the new method, a promoter is protected from degradation throughout the method. The promoter is constructed so that it does not serve as a primer for extension of a sequence that is complementary to the target sequence, i.e. it is blocked. The method can be combined with other processes to eliminate the need for multiple steps and varying reaction conditions and their associated problems. At least three otherwise separate enzymatic reactions can occur consecutively in one phase (i.e., without organic extraction and precipitation), more preferably in the same reaction extraction and precipitation), more preferably to the new method may occur in a modified low salt buffer. The present sequence represents the poly A tract of a mRNA used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 20
Query Match
Best Local
                                         Sequence 24
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                                                                                                                                                                                                                                                                                                                                            Amplification of nucleic acids, extension at the 3' end, useful
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                                                                                                                                                                                                                                                                                    The invention relates to a method of amplification of nucleic acid which
                                                                                                                                                                                                                                                                                                                Disclosure; Fig 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-MAY-2001; 2001US-00854317.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      varying reaction
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                                           BP; 24 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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/*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           synthesis primer"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amplification; multiple step elimination; condition elimination; poly A tract.
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0.3%;
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                                                                                                                                                                                                                                                                                                                   English.
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 Score 18.2;
Pred. No. 3.
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Pred. No. 3.6e+02;
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Matches 20
                                                                                                                                                                                                                                                                      The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The metad and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                                              Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                      ADB36806 standard; DNA; 24 BP
                                                                                                                                                                                                                                                    Sequence 24 BP; 0 A;
                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 18; 221pp; English.
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hypo-responsive subject; immunostimulatory.
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          Symthetic
                            hypo-responsive
                                       ds; allergy;
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                                                                                                  ADB36806;
                                                                                                                                                                                                                                                                                                                                                                                                                                  (BRAT/) BRATZLER R L. (PETE/) PETERSEN D M. (FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunostimulatory nucleic acid
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                                                          Immunostimulatory nucleic acid #420
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20; Conserv
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                             asthma; poly-G nucleic acid; aerosol formulation; ive subject; immunostimulatory.
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Pred. No. 3.6e+02;
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RESULT 198
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                             Bratzler RL,
                                                                                                                                                                                                         02-FEB-2001; 2001US-00776479
                                                                                                                                                                                                                                            08-MAY-2003.
                                                                                                                                                                                                                                                                            US2003087848-A1
                                                                                                                                                                                                                                                                                                                                         ds; allergy; asthma; poly-G nucleic acid; aerosol formulation;
hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                                                                                                                                                                                          04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADB37259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADB37259 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24 BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 11; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bratzler RL, Petersen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-FEB-2000; 2000US-0179991P
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                                               WPI; 2003-657977/62.
                                                                                                                                                                            03-FEB-2000; 2000US-0179991P
                                                                                                                                                                                                                                                                                                                                                                                          Immunostimulatory nucleic acid #873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BRAT/) BRATZLER R L. (PETE/) PETERSEN D M. (FOUR/) FOURON Y.
                                                                                                            (BRAT/) BRATZLER R L. (PETE/) PETERSEN D M. (FOUR/) FOURON Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
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Pred. No. 3.6e
0; Mismatches
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RESULT 199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Creating recombination products between two distinct nucleotide sequences, useful in the field of synthetic gene technology, and in segmenting a library, or a population or a collection of polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          recombination product; synthetic gene technology; butterfly; biliverdin binding protein; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADD31867 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method for creating a collection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 3; SEQ ID NO 106; 132pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-663477/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-JAN-2002; 2002US-00062188
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         госат
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Best Local Simi
Matches 20;
                        The invention is directed to novel methods of amplifying and detecting DNA using rolling circle amplification (RCA). The invention relates to detecting a target sequence (I), which involves using a capture probe (CP) that is complementary to an open circle probe and includes a cleavage site. The method comprises: attaching a capture probe (CP) that is complementary to an open circle probe and includes a cleavage site. The method comprises: attaching a capture probe (CP) to a occ cleavage site (CS), to form a device; treating CP with (I) and OCP for form a hybridisation complex (HC); treating CP with a ligase so that OCP is circularised, forming a second complex (HC2); treating CP with a cleavage agent, to cut at CS, and adding an extension enzyme (EE) and mucleotide triphosphates (NTPs) to form an extended CP, which is cleavage agent, to cut at CS, and adding an extended CP, which is cleavage agent, to the complex (HC2); treating (I) that comprises two target domains (TD1, TD2) and (I) that comprises two adjacent target domains. The method is used for detecting (I) that comprises two target agent, seed for detection, genotyping and/or quantification of target sequences, for research, clinical use, quality control or field testing, particularly detection of multiplexing, and since it provides that the control of multiplexing, and since it provides a second complex (HC2).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               open circle probe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting nucleic acid targets, useful e.g. for diagnosing single nucleotide polymorphisms, by extension of capture probe complementary to
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Pred. No. 3.
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                                                                                                                                                                                                                                                                                  The invention relates to methods for stimulating antibody dependent cellular cytotoxicity (ADCC), for modulating immune response and for inducing antigen-specific immune response which involve administering imidazoquinoline agents in conjunction with other agents. The method is useful for stimulating ADCC in a subject having a disorder chosen from asthma/allergy, infectious disease, cancer and warts. The present sequence is an immunostimulatory oligonucleotide used to illustrate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Stimulating antibody dependent cellular cytotoxicity, modulating immune response and inducing antigen-specific immune response in subject by administering imidazoquinoline agents in conjunction with other agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for direct detection and quantitation of unmodified targets. sequence is that of a single base extension (SBS) probe used genotyping with RCA signal amplification to demonstrate the minvention.
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   ss; non-CpG; immunostimulatory; non-palindromic; immune response;
proliferation; differentiation; cytokine; antibody production; B-
plasmacytoid dendritic cell; immunomodulator; gene therapy;
chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 18.2;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 24;
                                                                                                                                                             24.
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ADG7557
XX ADG7557
XX ADG7555
XX ADG7555
XX ADG7555
XX ADG755
XX A
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromnic sequence motif. Specifically, it refers to DNA oligonucleotides (without a Cpg motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune response is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is an immunostimulatory non-CpG carcinoma.
                                                                    WO2003101375-A2
                                                                                                                                                                                                                             88; CDG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cyrokine; antibody production; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG75917
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 24 BP; 1 A; 1 C; 1 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAY-2002; 2002CA-02388049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAY-2003; 2003WO-EP005691
                                                                                                                                                                                                                                                                                                                                                                                                 Non-CpG DNA oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           variant DNA oligo, used in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  renal cell carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (IMMU-) IMMUNOTECH SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5403
                                                                                                                                                                                                     cell carcinoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAAAAAGAAAAATGAAAATAAA 5425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAAAAAACAAAATGAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
ilarity 87.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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Pred. No. 3.6e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                 IMT 053 SeqID
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.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                 19.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
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Best Local S
                                                                                                                                                                                                                                              Matches 20;
                                                                                                                                                                                                                                                                                                                                                                      contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonuclectides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune and antibody production in B-cells, as well as cell differentiation, cytokine cytokine production in plasmacytoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing on ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonuclectide sequence is the non-CpG DNA oligo IMT carcinoma in a composition of the invention.
                                                                                                                                                                                                                                                                                                                     Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-MAY-2002; 2002CA-02388049.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAY-2003; 2003WO-EP005691.
                                                                          AAQ95960
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DN
 06-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 3; SEQ ID NO 19; 139pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (IMMU-) IMMUNOTECH SA
                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                        5393 АЛЛАЛАТАСАЛАЛАДАДАЛАЛА 5415
                                                                                                                                                                                                                                                                                                                                                              used
                                                                                                                                                                       24
                                                                          standard;
                                                                                                                                                                       AAAAAAAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                            in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                       BP; 0 A; 0 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                Conservative
(first entry)
                                                                          DNA;
                                                                                                                                                                                                                                                               0.3%;
                                                                          25
                                                                          BP
                                                                                                                                                                                                                                              Pred. No. 3.66
0; Mismatches
                                                                                                                                                                                                                                                                 Score 18.2;
Pred. No. 3.
                                                                                                                                                                                                                                                                   .6e+02
                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                       24;
                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The immune
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RESULT 205
AAQ959
AC AAQ959
AC AAQ959
AC AAQ959
XX Immobil
KW Immobil
KW Impbrid
KW Sequen
OS Synthe
XX Key
FT misc_f
FT misc_f
FT WO951!
XX WO951!
XX O6-DE(
XX O6-DE(
  06-DEC-1993;
                            06-DEC-1994;
                                                                                                                                                                                                                                                         Oligonucleotide biotin-T25 for novel nucleic acid immobilisation method
                                                     15-JUN-1995
                                                                               WO9515970-A1
                                                                                                                                  misc_feature
                                                                                                                                                                                                  Immobilisation; solid support; salt; cationic detergent; capture probe;
hybridisation; primer; template-dependent extension; target organism;
sequencing; genetic polymorphism; ss.
                            94WO-US014096.
                                                                                                                                                 Location/Qualifiers
                                                                                                          'note= "biotinylated
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RESULT 206
AXX84258/c
ID AAX842
XX AAX842
XX AX842
XX AX842
XX NCk as
KW therag
XX Synthe
OS Homo E
XX W09931
XX W09931
XX I1-DE(
XX I2-DE(
XX IX-DE(
XX IX-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotides AAQ95959-82 are examples of oligonucleotides used in a novel method of immobilising oligonucleotides to a solid support by incubating in the presence of a salt or cationic detergent e.g. NaCl (50-250 mM, pH 6.0-8.0) or 1-ethyl-3-(3'-dimethyl amino propyl)-1,3 carbodiimide hydrochloride (ECD). The oligonucleotides can be capture probes for detection of specific nucleic acids by hybridisation or can be primers for template-dependent extension from the immobilized primers on nucleic acid from a target organism. The method can be used in hybridisation assays, sequencing and analysis of genetic polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Immobilising synthetic nucleic acid on solid support - by incubation in presence of salt or cationic detergent, for use in hybridisation assays, sequencing and analysis of polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                    Sakaki Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-JUN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             W09931239-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nck associated protein 1; Napl; human; apoptosis; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer for human Nck associated protein 1 coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX84258;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX84258 standard; DNA; 25 BP
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                                                                                    Example 1; Page 76; 90pp; Japanese.
                                                                                                                                           Alzheimer's disease
                                                                                                                                                                  Protein inhibiting apoptosis, useful in
                                                                                                                                                                                                                             WPI; 1999-395181/33
                                                                                                                                                                                                                                                                                                                                                                                                                          15-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                        (KYOW ) KYOWA HAKKO KOGYO KK
(SAKA/) SAKAKI Y.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MOLE-) MOLECULAR TOOL INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5393 AAAAAAATACAAAAAAGAAAAAA 5415
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAAAAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 18; 61pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           94US-00341148
                                                                                                                                                                                                                                                                                                                                                                                                                          97JP-00363183
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 98WO-JP005646.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 18.2; DB 1;
Pred. No. 3.7e+02;
                                                                                                                                                                        the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                     diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                     and treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25;
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This sequence represents a PCR primer used to isolate DNA encoding the human Nck associated protein 1 (Napl) of the invention. Napl inhibits

RESULT 208
AAA39306/c
ID AAA393
XX
AC AAA393
XX
DT 11-SEP

AAA39306 standard;

RNA:

25 ₽P

11-SEP-2000 AAA39306;

(first entry)

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RESULT 207
AXX842560/c
ID AXX842
XX AAX842
XX AX8842
XX DOS SEP
XX PCR px
XX NCk as
KW Lherap
XX Synthe
OS Homo s
XX W09931
XX W99931
XX I1-DEC
XX I4-DEC
XX (KYOW
PA (SAKA)
PI Sakaki
XX WPI;
PI Sakaki
XX Protei
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XX This (CC human
CC apopt(
CCC treati
XX Sequei
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                                                              Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local (
                                                   Matches
                                                                                                                          This sequence represents a PCR primer used to isolate DNA encoding the human NCk associated protein 1 (Napl) of the invention. Napl inhibits apoptosis. The protein can be used in the investigation, diagnosis and treatment (e.g. by gene therapy) of Alzheimer's disease
                                                                                                    Sequence 25 BP; 0 A; 1 C; 0 G; 24 T; 0 U; 0 Other;
                                                                                                                                                                                              Disclosure; Page 77; 90pp; Japanese.
                                                                                                                                                                                                                                                              WPI; 1999-395181/33
                                                                                                                                                                                                                                                                                      Sakaki Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nck associated protein 1; Napl; human; apoptosis; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primer for human Nck associated protein 1 coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX84260;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX84260 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  apoptosis. The protein can be used in the investigation, treatment (e.g. by gene therapy) of Alzheimer's disease
                                                                                                                                                                                                                        Alzheimer's disease
                                                                                                                                                                                                                                   Protein inhibiting apoptosis, useful in the diagnosis and treatment of
                                                                                                                                                                                                                                                                                                                                                        15-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                               14-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                         24-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9931239-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapy; PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                  (KYOW ) KYOWA HAKKO KOGYO KK
(SAKA/) SAKAKI Y.
                        5393 ААААААТАСААААААСААААА 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5393 AAAAAAATACAAAAAAGAAAAAA 5415
24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20;
                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25 BP; 0 A; 0 C; 1 G; 24 T; 0 U;
AAAAAAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer; 88.
                                                                                                                                                                                                                                                                                                                                                        97JP-00363183
                                                                                                                                                                                                                                                                                                                                                                                  98WO-JP005646
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.3%;
87.0%;
                                                             0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP.
                                                   <u>.</u>.
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                                                              Score 18.2; DB 1
Pred. No. 3.7e+02
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Pred. No. 3
                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .7e+02
                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 Other;
                                                                         Length 25;
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                                                   Indels
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                                                   Gaps
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RESULT 209
AAZ30267/c
ID AAZ302
XX AAZ302
XX AAZ302
XX Captur
XX Captur
XX C-myc
KW C-myc
KW nuclei
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                                                                                                                                                                                                                                                                                                                                                                            δ
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a rapid pairing reagent (1) for the isolation or detection of a polymucleotide (PN) analyte molecule having selected target base sequence, in a sample containing the analyte molecule and non-target polymucleotide, comprising a capture component (A) and a target-specific probe (B) linked to a solid substrate. The
                                                                                                                c-myc fusion; non-nucleoside spacer; capture probe;
nucleic acid-protein fusion; ribosome display parti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probes are not extended by polymerases and therefore do not interfere with amplification of target molecule. AAA39306 to AAA39316 represent oligonucleotides used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            isolated sequences are useful for enzymatic amplification. (I) is capable of rapidly binding nucleic acids in the sample and placing them in close proximity to target probes on the reagent, thus enabling binding under low stringency. Combination of rapid capture and concentration of polymucleotides with selective targeting of analyte molecules, greatly enhances the isolation process. Non-ionic morpholino oligomers used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reagent for rapidly detecting or isolating target nucleic acid sequences in polynucleotide-containing sample, comprises capture component and target-specific probe linked to solid substrate.
                  14-OCT-1999
                                                                                 Synthetic
                                                                                                                                                                Capture probe CP125 specific for c-myc fusion targets
                                                                                                                                                                                                  11-FEB-2000
                                                                                                                                                                                                                                 AAZ30267;
                                                                                                                                                                                                                                                                 AAZ30267 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25 BP; 0 A; 0 C; 0 G; 0 T; 25 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; Col 17; 24pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-364413/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Wages JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rapid detection; probe; target nucleic acid; enzymatic amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rapid
                                                   WO9951773-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AVIB-) AVI BIOPHARMA INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          solation;
                                                                                                                                                                                                                                                                                                                                                                                  5393 AAAAAAATACAAAAAAGAAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          capture probe designated Neu-probe SEQ ID NO:1.
                                                                                                                                                                                                                                                                                                                                                 25
                                                                                                                                                                                                                                                                                                                                                 AAAAAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Summerton JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             96US-0030963P
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                                                                                                                                                                                                                                                                                                                                                                                                                              0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Weller DD;
                                                                                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                              Score 18.2; DB 1;
Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                              display particle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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The present sequence represents a capture probe specific for a c-myc (fusion target. It is used in the method of the invention. The specification describes the use of non-nucleoside spacers to immobilise a probes of capture probes on a solid support. The solid support carries an array of capture probes, each consisting of non-nucleoside spacers consisting of non-nucleoside spacers prevent a ribosome display particle is bound. Non-nucleoside spacers prevent interaction of proteins with the support surface, ensuring efficient hybridiation between capture probes and bound nucleic acid/protein fusions, while mainimising denaturation of the protein which may then consisting, its native folded structure. The arrays of capture probes are used to screen for interactions between proteins and compounds (e.g. other proteins, ligands or nucleic acids), particularly to identify potential therapeutic agents, enzyme substrates or unknown proteins that interact with drugs, but also for diagnosis (detecting disease-associated with drugs, but also for diagnosis (detecting sin a sample
  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Attaching capture probes to solid phases through non-nucleic spacers, producing arrays for detecting interactions of proteins with other compounds, e.g. for drug screening.
                                                                    Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 8; Page 29; 57pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PHYL-) PHYLOS
                                                                                                                 and for quantifying target molecules in a sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wagner
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0.3%;
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ABK499
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ABK499
XY
AC
ABK499
DT
15-JUL
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Glass-
KW
Glass-
KW
Chroma
XX
Synth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chromatography
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Glass-synthetic resin membrane; pore glass-polytetrafluoroethylene resin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example oligonucleotide #2 prepared on glass-synthetic resin membrane.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABK49986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK49986 standard; DNA; 25
                                                                                                                        Wong YN,
                                                                                                                                                                                                                                                                                                21-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                             04-MAY-1999;
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                                                                                                                                                                                                              (CPGC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5393 AAAAAAATACAAAAAAGAAAAA 5415
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                                                                                                                                                                                                              CPG INC.
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                                                                                                                                 Chen R;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                96US-00604440
                                                                                                                                                                                                                                                                                                                                                                                 99US-00305219.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                membrane; PTFE; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 18.2; DB 1; Length 25; Pred. No. 3.7e+02; O; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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WPI; 2001-534961/59

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ARBSULT 211
ABV80873
ID ABV808
XX ABV808
XX ABV808
XX ABV808
XX Human
XX Human
XX Human
XX Human
XX Human
XX Prosta

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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                  30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-MAY-2001;
09-OCT-2001;
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                            Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and for treating subjects having defects in HTPL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kithey; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Preparation of controlled pore glass-polytetrafluoroethylene resin chromatography membrane by heating, calendering and sintering mixture of controlled pore glass and aqueous dispersion of polytetrafluoroethylene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25 BP; 0 A; 0 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-JAN-2002; 2002EP-00001167.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
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                                                                                                                                                       2002-676582/73
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                                                                                                                                                                                                                                                                            AEOMICA
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2001WO-US000664.
2001WO-US000667.
2001WO-US000667.
2001WO-US000668.
2001WO-US000668.
2001WO-US000669.
2001US-00864761.
2001US-0327898P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                               INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 18.2; DB 1;
Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ij
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ABSULT 212
ABV8084
ABV8084
AC ABV808
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CC has two isoforms, with a few single base pair differences between the
CC two. One of the single base pair changes introduces a premature stop
CC codon in HTPL-S (S for short) compared to HTPL-U (L for long). HTPL
CC shares an overall structure organisation with the Patched protein. The
CC shared structural features strongly imply that HTPL plays a role similar
CC to that of Patched, and is a potential tumour suppressor. HTPL is
CC important in regulating male germ cell development, and the HTPL gene was
CC mapped to human chromosome 10p12.1. HTPL and its coding sequence are
CC useful for diagnosing a disorder caused by mutation in HTPL, and in
CC therapy and manufacture of a medicament for treatment or prevention of
CC such disorder associated with decreased expression or activity of human
CC HTPL. Such disorders include disorders of testis, or adrenal, adult and
CC foetal liver, bone marrow, brain, kidney, lung, placenta, prostate,
CC skeletal muscle or colon function. HTPL proteins and nucleic acids are
CC clinically useful diagnostic markers and potental therapeutic agents for
CC male infertility and cancer. The present oligonucleotide was used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                             30-JAN-2001; 2001WO-US000663.
30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000665.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001WO-US000669.
23-MAY-2001; 2001US-00864761.
09-OCT-2001; 2001US-0327898P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
                                                                                            WPI; 2002-676582/73.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-JAN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human HTPL scanning oligonucleotide SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABV80874 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 25
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                                                                                                                                                                                                                         AEOMICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCCCAAGCCCAGGCCGGGGCCGG 24
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Pred. No. 3
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Novel isolated human testis expressed Patched like protein (HTPL), useful for identifying agonist and antagonist and specific binding partners, and

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                            Human connexin 9 and encoded polynucleotide, treatment of malignant tumor, hemopathy, HIV diseases and inflammation.
                                                                                                                                                                                                                                                                                                                           WO200181538-A2.
                                                                                                                                                                                                                                                                                                                                                                                                   Human; connexin 9; cytostatic; virucidal; immunomodulatory; antiinflammatory; haemostatic; malignant tumour; haemopathy;
                                                 Example
                                                                                                                                          WPI; 2002-034440/04.
                                                                                                                                                                                                                                 27-APR-2000; 2000CN-00115456
                                                                                                                                                                                                                                                               23-APR-2001; 2001WO-CN000608
                                                                                                                                                                                                                                                                                               01-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human connexin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        example from the invention
                                                                                                                                                                                                    (BIOW-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABA03917 standard;
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                                                                                                                                                                                                      BIOWINDOW GENE DEV INC SHANGHAI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page
                                               Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 9 PCR primer 2
                                                                                                                                                                                                                                                                                                                                                                                    immunological disease; inflammation; PCR primer;
                                              12; 32pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 25; .7e+02;
                                                                                          applicable in diagnosis and infection, immunological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
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The present invention describes human connexin 9 virucidal, immunomodulatory, antiinflammatory and

(I). (I) has l haemostatic

cytostatic, activities.

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RESULT 214
ACK28658
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                                                                                                                                                           The invention discloses a microarray comprising a plurality of nucleic caid probes including one of 2,018,500 fully defined sequences, or its comperfect match, perfect mismatch, antisense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, compounds. The nucleic acid probes are specifically designed for analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises comprises and detecting the hybridisation. The nucleic acid probes are antached to a solid support. The analysis comprises monitoring gene expression levels, identifying biallelic markers or polymorphisms, cor family members of a gene and a cross-species comparison. Each of the nucleic acids further comprises a tag sequence. The array of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dotter the accounter of the contribution of the contribution of the contribution of the contribution.
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Best Local Simi
Matches 20;
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blot hybridisation to identify or detect the sequence or specific mutations of any gene, in mapping the 5 termini of mENA molecules by primer extensions or in screening cDNA or genomic libraries or subclones for additional subclones containing segments of DNA that have been isolated and previously sequenced. The sequence presented is one of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACK28658 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New array of nucleic acid probes, useful for in Southern, Northern or dot-blot hybridization to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-567953/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-MAR-2002; 2002US-00098263.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human microarray DNA oligonucleotide SEQ ID NO 128639.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-MAR-2001; 2001US-0276759P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or specific mutations
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Pred. No. 3.
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RESULT 215
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Probe carrier manufacturing method for inkjet liquid discharge head in direction orthogonal angle satisfying predetermined relation.
                      Oligonucleotide
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                                                                  18-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nucleic acid probes incorporated in the microarray. Note: The sequence data for this patent can also be obtained in electronic format directly from USPTO at segdata.uspto.goc/sequence.html
                                                                                                                                                   ADC54008 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      28-MAR-2001; 2001JP-00094400
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Similarity 87.0%;
20; Conservative
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                    of the invention
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                                                                                                                                                     DNA;
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Pred. No. 3.7e+02;
0; Mismatches 3;
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Pred. No. 3.
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to scanning direction, at
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RESULT 217
ADF39737/c
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XX Probe
XX Probe
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XX Z8-MAJ
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Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Probe array; microarray; DNA array; DNA chip; manufacture; electrostatic adsorption mechanism; DNA analysis; simultaneous gene detection; probe; ss.
                                                                                                                             Manufacturing of probe carrier for carrying probes for base sequence analysis of genetic deoxyribonucleic acid and simultaneous multiple diagnosis of gene by ink jet process while removing mist of probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 25 BP; 25 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-JAN-2003
                                                                                                                                                                                                                                                                                                                                                  (CANO ) CANON KK
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Pred. No. 3.7e+02
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Matches 20
                                                                  Query Match
                                                                                                                                                                                          The invention relates to a method and device for the manufacture of a probe array. The method involves using an inkjet system to discharge a probe solution through a solution discharging head, so as to form a number of probes on a solid matrix. Mists of the probe solution generated during probe solution discharge are caught by an electrostatic adsorption mechanism. The method and device are suitable for manufacturing probe arrays for analysing DNA sequences, and for the simultaneous detection of multiple genes. The method and device of the invention prevent the scattering of probe positions and the mixing of different probe solutions. The present sequence is related to the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Manufacturing of probe carrier for carrying probes for base sequence analysis of genetic deoxyribonucleic acid and simultaneous multiple item diagnosis of gene by ink jet process while removing mist of probe
                                                                                                                                     Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 2; SEQ ID NO 3; 15pp; Japanese.
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Score 18.2; D
Pred. No. 3.7e
0; Mismatches
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Pred. No. 3
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                                                                                                                                                                                              cc or more polymorphic microsatellite loci (PML). The method comprises: PCR campilfication of at least one DNA region of (I) that includes PML, using CC as template a DNA sample containing at least one segment of (I); and CC determining the length of the resulting amplicon(s). Also described are: a method of determining (M2) microsatellite markers (MM) for CC predisposition to a disease, associated with a gene that includes one or CC more PML; and prediagnosis (M3) of diseases associated with gene that include or CC include PML. The method is used to identify microsatellite markers, in a CC disease related gene, that are associated with a predisposition to CC diseases and for prediagnosis of such diseases, especially prion diseases CC but also cystic fibrosis, malignant hyperthermia syndrome in pigs and CC metabolic diseases; also to type genes that encode milk proteins, CC hormones or transcription factors. The method is simpler, quicker and CC particularly less expensive than known methods based on sequencing. This cc consensus seminores a prion protein polymorphic microsatellite marker CC consensus seminores.
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Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic disease; milk protein; hormone; transcription factor; pT7-blue-vector; sheep;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADO81145
                                                                                                                                                                                     consensus sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention describes a method of typing (M1) a gene (I) that has one
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-AUG-2002; 2002DE-01036711.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prion protein polymorphic microsatellite marker consensus sequence #23
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                                                                       Local Similarity
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                                                                                                                                             BP; 0
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                                                                                                                                             G; 25 T; 0 U;
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Pred. No. 3.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention relates to a novel mini-sequencing type DNA coligonucleotide chip. Specifically, it refers to a chip that is useful for detecting methylation of promoter CpG islands occurring in multiple CC genes. The present invention describes using oligonucleotide primers to constitutes treating DNA of the target gene with sodium bisulfite in CC constitutes treating DNA of the target gene with sodium bisulfite in CC constitutes treating DNA of the target gene with sodium bisulfite in CC constitutes treating DNA of the target gene with sodium bisulfite in CC confirm the hypomethylation specific (MSP) PCR or multiplex MSP PCR to confirm the hypomethylation site of the promoter CpG islands of CC multiple genes. Accordingly, the chip comprises primer sequences designed CC from these PCR products that have amine linkers of 12 carbons attached to CC the 5'-terminal, which are spotted onto the glass slide coated with 3-cc aminopropyltrimethoxylan and 1,4-dissothiocynnate using an array robot. CC minopropyltrimethoxylan and 1,4-dissothiocynnate using an array robot. CC accurately and rapidly detecting methylation of CpG islands of multiple GC accurately and rapidly detecting methylation of CpG islands of multiple CC genes. This oligonucleotide sequence is a PCR primer used on the biochip, Given in an exemplification of the invention.
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Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 25 BP; 18 A; 4 C; 0 G; 3 T; 0 U; 0 Other;
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                                                                                                         09-SEP-2004 (first entry)
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                                   PCR primer amplifies promoter CpG islands of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
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Pred. No. 3.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                          cancer related
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RESULT 222
AAN70276/c
ID AAN702
XX AAN702
XX 03-OCT
DT 26-MAY
XX Sequen
XX Sequen
XX Hybrid
XX Synthe
XX Synthe
PN EP2279
XX EP2279
XX O8-JUI
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Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       amplify the sodium bisulfite treated DNA and sequencing the PCR product to confirm the hypomethylation site of the promoter CpG islands of multiple genes. Accordingly, the chip comprises primer sequences designed from these PCR products that have amine linkers of 12 carbons attached to the 5'-terminal, which are spotted onto the glass slide coated with 3-aminopropyltrimethoxylan and 1.4-dissothlocyanate using an array robot. The resulting minisequencing dissothlocyanate using an array robot. The resulting minisequencing dissothlocyanate using an array robot. The resulting minisequencing dissothlocyanate using an array robot. The rosulting side detecting methylation of CpG islands of multiple genes. This oligonucleotide sequence is a PCR primer used on the biochip, given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mini-sequencing; CpG island; methylation specific PCR; MSP; multiplex MSP PCR; cancer; PCR; primer; 88; microarray chip
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to a novel mini-sequencing type DNA oligonuclectide chip. Specifically, it refers to a chip that is useful for detecting methylation of promoter CpG islands occurring in multiple genes. The present invention describes using oligonucleotide primers to determine the position of a target gene and promoter CpG islands, this constitutes treating DNA of the target gene with sodium bisulfite in order to carry out methylation specific (MSP) PCR or multiplex MSP PCR to
                                                                                                                                                                                                             03-OCT-2002
26-MAY-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Minisequencing type oligonucleotide chip for detecting methylation of promoter CpG islands of multiple genes, useful for detecting cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-MAY-2002; 2002KR-00025108
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                                   08-JUL-1987
                                                                                                      Synthetic
                                                                                                                                      Hybridisation,
                                                                                                                                                                         Sequence of scissile link probe MRC060 (HL).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 573; 248pp; Korean
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-095256/10
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(first entry)
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Pred. No. 3.7e+02;
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04-DEC-1986;

86EP-00116906.

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RESULT 223
AAN70275/c
ID AAN702
XX AAN702
XX AAN702
XX Gequen
XX Sequen
XX Hybrid
XX Synthe
XX Synthe
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XX O8-JUI
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Matches 20
                                                                                                                                          Synthetic linked by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The patent claims a new molecule of formula (NA1---S---NA2)n. NA1 and NA2 are noncomplementary nucleic acid sequences; ---S--- a scissile linkage; n= 1 or 1,000, which is used for the detection of specific DNA or RNA sequences in a test soln. The scissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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The patent claims a new molecule NA2 are noncomplementary nucleic linkage; n= 1 or 1,000, which is
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26-MAY-1991
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                                                                                                                                                                                                                                                                                                     MEIOGENICS INC.
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                                                                                                                                        nucleic acid probes a scissile linkage.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       scissile link probe MRC059 (HL).
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Pred. No. 3
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  of formula (NA1----S----NA2)n. acid sequences; ---S--- = a sc: used for the detection of spec:
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31-OCT-2002
25-APR-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            or RNA sequences in a test soln. The scissile link probes may be PL (Permanent Linkage to Solid Support) or HL (Hydrolysable Linkage to Solid Support). The differential liability of DNA and RNA may be exploited in a heterogenous system when the scissile linkage is an RNA molecule. In the examples, counter probe molecules 9 through 16 were used to determine suitable hybridisation conditions. (Updated on 03-OCT-2002 to add missing OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                AAN92241 standard;
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                                                                                                                                                                                                                                                                                       misc_feature
                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                           Probe MRCO59; solid support; ribonuclease.
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                                                                                                                                                          (MBIO-)
                                                                                                                                                                           29-APR-1988;
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                                                                                                                                                          MEIOGENICS INC
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(revised)
(first en
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                                                                                                                                                                                                                                                                       /note=
                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                              note= "deoxyribonucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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Pred. No. 3
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probe MRCO59 is bound by a hydrolysable linkage to a solid support at its 3' end. It is used by reacting excess probe with a target nucleic acid; nicking hybridised probe at least new within a predetermined sequence to form 2 or more probe fragments hybridised to the target sequence, which results in the probe fragments becoming hybridised to another probe; and identifying probe fragments, so detecting the target sequence. The probe can react with target sequence to complete a cycling sequence. Using this

sensitivity of

of 10 exp. -19

exp.

10

exp.

Disclosure; Page 24; 34pp;

English

Detecting target nucleic acid molecules - using excess complementary nucleic acid probes and nicking to complete a cycling sequence.

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RESULT 225
AAN92242/c
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31-OCT-2002
25-APR-1990
        probe MRCO60 is bound by a hydrolysable linkage to a solid support at its 3' end. It is used by reacting excess probe with a target nucleic acid; nicking hybridised probe at least once within a predetermined sequence to form 2 or more probe fragments hybridised to the target sequence, which results in the probe fragments becoming hybridised to another probe; and identifying probe fragments, so detecting the target sequence. The probe can react with target sequence to complete a cycling sequence. Using this system, sensitivity of 10 exp. -19 to 10 exp. -20 molecules of target can be obtd. The probe is cleavable at the ribonucleotides by a ds RNase, eg RNase H or ExoII. (Updated on 31-OCT-2002 to add missing OS field.) (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                                            Detecting target nucleic acid molecules - using excess complementary nucleic acid probes and nicking to complete a cycling sequence.
                                                                                                                                                                                                                                                              Duck P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SS probe MRCO60.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAN92242 standard; DNA; 26 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 be obtd. The probe is cleavable at the ribonucleotides by RNase H or ExoIII. (Updated on 31-OCT-2002 to add missing (Updated on 25-MAR-2003 to correct PR field.)
                                                                                                                                                                   Disclosure; Page 24; 34pp; English.
                                                                                                                                                                                                                                                                                                                  29-APR-1988;
                                                                                                                                                                                                                                                                                                                                           29-APR-1988;
                                                                                                                                                                                                                                                                                                                                                                        02-NOV-1989
                                                                                                                                                                                                                                                                                                                                                                                                WO8910415-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Probe MRCO60; solid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAN92242;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 26 BP; 0
                                                                                                                                                                                                                                                                                        (MEIO-) MEIOGENICS
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                                                                                                                                                                                                                                      1989-339977/46.
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20; Conservative
                                                                                                                                                                                                                                                              Bender R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (revised)
(revised)
(first en
                                                                                                                                                                                                                                                                                                                  88US-00187814
                                                                                                                                                                                                                                                                                                                                            88US-00187814
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "ribonucleotides."
17. 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       note= "deoxyribonucleotides."
                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "deoxyribonucleotides."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 0 C; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 18.2; DB 1;
Pred. No. 3.7e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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OS field.)
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RESULT 226
ANT93819/c
ID ANT938
XX ANT938
XX ANT938
XX PAOSP
ANT 25-MAR
DT 24-FEB
XX Phosph
XX Phosph
XX Phosph
XX Iymphc
OS Synthe
CC NOVEL
CC IXMPI;
IYM WPI;
IYM New pl
PT Ne
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                                                               Novel phosphodiesteric oligonucleotides AAT93811-27 are based on the generic formula, in the 3'-5' or 5'-3' direction: (GaTa')a''-(GbTb')b''-(GGTc')c''-(GdTc')c''-(GGTc')c''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(GfTf)f''-(G
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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                liquid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New phospho:di:esteric oligo:nucleotide(s) - which exert a specific and selective cytotoxic effect on tumour cells, for treating both solid and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1997-319771/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scaggiante B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-DEC-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antitumoural phosphodiester oligonucleotide 9 with cytotoxic activity
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24-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT93819,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-JUN-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (SAIC-) SAICOM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 5; 38pp; English.
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "phosphodiester oligonucleotide"
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Pred. No. 3.
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Query Match

Sequence 26

BP; 0 A; 0 C; 2

G; 24 T; 0 U;

0 Other;

0.3%;

Score 18.2;

DB 1; Length 26;

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RESULT 228
AAX78723/c
ID AAX787
XX
AC AAX787
XX
DT 03-SEP
XX
DT 03-SEP
XX
DB Human
XX
KW Pancre
KW anti-P
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AAX07466/C
ID AAX074
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity Matches 20; Conser
                                                                                                                                                                                                                                                                                                                                                             Billing-Medel PA, Cohen M,
Granados EN, Hodges SC, K
Scheffel CP, Stroupe SD,
                                                                                                                                                                                                                                   The sequence is that of an oligonucleotide used in the isolation of BS124-specific EST clone. It is useful for detecting, diagnosing, staging, preventing or treating, or determining predisposition to diseases or conditions of the breast, such as breast cancer
Pancreatic disease; PA153; human; cytostatic; anti-PA153; antagonist; therapy; treatment; to
                                                                                                                                                                                                                 Sequence 26 BP; 0 A; 0 C; 1 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                     New isolated BS124 polynucleotides diagnosing, preventing or treating such as breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-JUN-1999
                              Human pancreatic PA153 EST-specific clone primer 12.
                                                  03-SEP-1999
                                                                                       AAX78723 standard;
                                                                                                                                                                                                                                                                                   Disclosure; Page 97; 125pp; English.
                                                                                                                                                                                                                                                                                                                                             WPI; 1999-105623/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                           19-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BS124; breast; cancer; detection; diagnosis; prevention; treatment; EST;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human BS124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX07466 standard; cDNA; 26 BP.
                                                                                                                                                                                                                                                                                                                                                                                                     (ABBO ) ABBOTT LAB.
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                                                                                                                                                         5393 АЛАЛАЛАТАСАЛАЛАДАДАЛАЛА 5415
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                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                      97US-00879354
                                                                                                                                                                                                                                                                                                                                                                                                                                           98WO-US012862
                                                                                       DNA;
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Yu H;
                                                                                       ВP
                                                                                                                                                                           0,
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                                                                                                                                                                                    Score 18.2; DB 1;
Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                  Colpitts TL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 3.7e+02;
0; Mismatches 3;
                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                               and polypeptides - used for detecting, diseases or conditions of the breast,
                                                                                                                                                                                                                                                                                                                                                                         Kratochvil JD,
                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                         Friedman PN, Gordon J; hvil JD, Russell JC;
  c; detection; antigen;
tumour; metastasis;
                                                                                                                                                                                             Length 26
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  RESULT 229
AAF77536/c
ID AAF775
XX AAF775
XX AAF775
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XX CDNA 1
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XX CDNA 1
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XX PR 20-NON
XX CCHUO)
PA (CHUO)
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Best Local S
Matches 20
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                                                                                      US6197554-B1
                                                                                                         Synthetic.
                                                                                                                                                                                                        AAF77536;
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  (LINS/) LIN S. (CHUO/) CHUONG
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This invention describes novel contiguous and partially overlapping cDNA sequences and their encoded polypeptides, designated PA153, transcribed from human pancreatic tissue and which have cytostatic activity. The PA153 polynucleotides, proteins and antibodies are all useful in methods of detection. Detection of PA153 polynucleotide, antigens or anti-PA153 antibodies in a sample is indicative of pancreatic disease. PA153 antibodies (antagonists) can also be used in vivo for therapeutic use, e.g. treatment of pancreatic disease, tumours or metastases. Antisense PA153 polynucleotides can be used in gene therapy of pancreatic diseases. AAX78712-X78725 represent primers used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Billing-Medel PA, Cohen M, Colpitts TL, Granados EN, Hodges SC, Klass MR, Kratt Russell JC, Stroupe SD;
                                                                                                                                                                                                                                                                             cDNA library production; SCLA; gene chip technology; differential screening; pathological diagnosis; gene single-cell cDNA library amplification; ds.
                                                                                                                                                                                                                                                                                                                                                                                            CDNA library production method related oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAF77536 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PA153 cDNA transcribed from pancreatic tissue
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98US-00197951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97US-00990568
                                                       98US-00197951.
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Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Friedman PN, Gordon J;
chvil JD, Roberts-Rapp
                                                                                                                                                                                                                                                                                                          genetic identification,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
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ARESULT 230
AAP23526/c
ID AAP235
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AC AAP235
XX Primer
XX Uniden
XX Uniden
XX (ALNS/PP 04-JUN
PP 04-JUN
PP 04-JUN
PR (CHUO/PA (WIDE/XX
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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                    (LINS/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Generating a complete full-length cDNA library from single cells for in gene chip technology, involves reverse transcribing intracellular mRNAs, adding polynucleotide tail and amplifying formed cDNAs.
                 The present invention relates to generating amplified messenger RNAs with polymerase reaction activity, comprising cycling steps of reverse transcription, denaturation, double-stranded cDNA synthesis and in vitro transcription. The invention is used for generating amplified mRNAs from limited mRNAs from single cells
                                                                                                                                                                Generating amplified messenger RNA sequences from single cells, involves cycling steps of reverse transcription, denaturation, double-stranded DNA sequences and in vitro transcription.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primer #4.
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                                                                                                                               Disclosure, Page 17; 31pp; English.
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                                                                                                                                                                                                                                                                              Lin S,
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') YING S.
') CHUONG C.
') WIDELITZ R B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mRNA;
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87.0%;
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                                                                                                                                                                                                                                                                              Widelitz
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Pred. No. 3
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Best Local S
Matches 20
                                                                                                                                                                                                                                                            This sequence represents a molecular scaffold which may be used in a composition of matter comprising multiple fluorophores. The fluorophores are bound to the molecular scaffold at separate predetermined positions, to permit fluorescence energy transfer between two fluorophores. The fluorophores are characterized by maximum emission wavelength of one being greater than the minimum excitation wavelength of the other. The composition is useful for determining whether a preselected nucleotide residue is present at a predetermining whether a preselected nucleotide is also useful in multicomponent analysis including multiplex biological analysis, and identifying multiple single nucleotide polymorphisms. The presence of a number of given nucleotide residues is determined simultaneously by the composition of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Composition of matter useful for multi-component analyses, comprises multiple fluorophores bound to molecular scaffold at preset positions to permit fluorescence energy transfer between two fluorophores.
                                                                                                                                                                                                                  Sequence 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 43; 113pp;
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31-JUL-2001; 2001US-0309156P.
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                                                                                                                                     Similarity
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                                                     AAAAAAATACAAAAAAGAAAAAA 5415
                                                                                                                                                                                                                  BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; 0 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
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                                                                                                                                  0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26
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                                                                                                                                     Score
Pred.
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Pred. No. 3.
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                                                                                                           Mismatches
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                                                                                                                                     18.2;
No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .7e+02
                                                                                                                                                                 DB
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                                                                                                                                                            Length
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                                                                                                                                                                 26;
                                                                                                           0,
                                                                                                           Gaps
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                                                                                                              0
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AAS206772/c
IID AAS206
XX AAS206
XX AAS206
XX Oy-AAPR
XX Human
XX Cytoki
KW Gytoki
PF 09-MAI
XX O9-MAI
XX O9-MAI
XX O9-MAI
XX IXMO
XX IXMO
XX IXMO
XX New Z
FT Gross
XX WPI;
XX New Z
FT Stimul
PT Gross
XX WPI;
XX New Z
FT Stimul
XX The p
CC Zalph
XX Gross
XX The p
CC Zalph
CC Ganti-
PT of inl
PT 
RESULT 233
AAD43853/c
ID AAD438
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to the isolation of a novel cytokine, CC zalphall Ligand and the polynucleotide encoding it. The invention also CC gives the sequence for the zalphall receptor and the polynucleotide CC encoding it. The zalphall Ligand polypeptide stimulates proliferation of CC natural killer (NK) cells or NK cell progenitors, the activation of NK CC cells, proliferation of T-cells, proliferation of B-cells stimulated with CC anti-CD40 antibodies, stimulates an antigenic response in a mammal, and CC reduces proliferation of B-cells stimulated with anti-IgM antibodies. The CC zalphall Ligand polypeptide is also useful in preparing antibodies that CC bind to zalphall Ligand epitopes. The zalphall Ligand polypucleotides can CC used as probes or primers to clone regions of a zalphall Ligand gene, CC and in gene therapy. Zalphall Ligand may also be used to identify CC inhibitors of its activity, to enhance the generation of anti-tumour CC responses with or without the infusion of donor lymphocytes, and to activate or stimulate the immune system. The present sequence represents the sequencing primer used to sequence CDNA clones in the isolation of the control of the
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 20
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11-MAR-1999;
01-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New zalphall ligand polypeptides and polynucleotides, useful for stimulating proliferation, activation, differentiation and/or induction of inhibition of specialized cell function, or for stimulating an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novak JE,
Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cytokine; zalphall Ligand; zalphall receptor; NK cell progenitor; natural killer cell proliferation; T-cell proliferation; B-cell proliferation; anti-tumour response; immune system;
                                       AAD43853 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 7; Col 139; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-040208/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human zalphall Ligand sequencing primer ZC7764b.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAS20672;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antigenic response.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ZYMO ) ZYMOGENETICS INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunostimulant; cytostatic; human; sequencing primer; ss.
                                                                                                                                                                                                                                                                                                      5393 AAAAAAATACAAAAAGAAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26
                                                                                                                                                                                                                            АЛАЛАЛАЛАЛАЛАЛАЛАЛА 3
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Johnston JV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 0
                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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99US-0123904P.
99US-0142013P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; 1 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                 0.3%;
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                                           ₽P
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                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                             Score 18.2; DB 1
Pred. No. 3.7e+02
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 26;
                                                                                                                                                                                                                                                                                                                                                                                     Indels
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RESULT 234
ABX93461/c
ID ABX934
XX ABX934
AC ABX934
XY 27-MAY
XX LS147-
XX LS147;
KW LS147;
KW primer
XX X
CS Synthe
XX
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                                                                                                                                                                                                                                                                                                             S
                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a method for obtaining a single stranded polynucleotide tag from a biological sample by cleaving one of the complementary strands of a double stranded polynucleotide with a cleavage agent capable of recognising a double stranded polynucleotide comprising complementary strands and cleaving only one of the strands of the polynucleotide in the process of generating a single strands of polynucleotide in the process of generating a single stranded polynucleotide tag. The method is useful for separating, analysing tags originating partly, and preferably wholly from a source of DNA and/or RNA in a sample comprising biological cells. The method is particularly for analysing gene expression (expression profiling or differential gene expression, or in diagnosing clinical conditions. The present sequence is a primer used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 26 BP; 0 A; 0 C; 0 G; 26 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Obtaining single stranded polynucleotide tags from a biological sample, for analyzing gene expression or diagnosing clinical conditions, comprises employing nicking endonucleases that cleave complementary
                                                 LS147; cancer; lung cancer; gene therapy; cytostatic; ss; primer; EST clone; expressed sequence tag clone.
                                                                                                  LS147-specific
                                                                                                                                   27-MAY-2003
                                                                                                                                                                   ABX93461;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-JAN-2001; 2001DK-00000126.
12-FEB-2001; 2001US-0267704P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Single stranded polynucleotide tag; cleavage agent; gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Primer #2 used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAD43853;
                                                                                                                                                                                                  ABX93461 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example; Page 294; 302pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-636542/68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (GENO-) GENOMIC EXPRESSION APS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-JAN-2002; 2002WO-DK000052.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-AUG-2002
                                                                                                                                                                                                                                                                                                                  5393 AAAAAAATACAAAAAAGAAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                 polynucleotide sequencing related universal primer #1.
                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                              0.3%;
                                                                                                                                                                                                   26
                                                                                                                                                                                                   ΒP
                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                Score 18.2; DB 1;
Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                   0 Other;
                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 26;
                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                    sequencing;
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RESULT 235
ABZ24784/C
ID ABZ2477
XX ABZ2477
XX O7-APR
XX Immuno
XX Immuno
XX Immuno
XX Synthe
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            맑
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention describes a purified polypeptide or its fragment derived from the LS147 gene capable of selectively hybridising to the nucleic acid of the gene and has at least 50% identity with the polynucleotide. The LS147 polypeptide is useful for preparing a composition for treating cancer, e.g. lung cancer using gene therapy. This sequence represents a universal primer used to sequence LS147 expressed sequence tag (EST)-
                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligodeoxynucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-APR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ24784 standard; DNA; 26 BP
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Russell JC,
21-MAY-2001; 2001AT-00000805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ24784
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 39; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BILL/)
(COHE/)
(COLP/)
(FRIE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUN-1998;
                                                17-MAY-2002; 2002WO-EP005448
                                                                                                                                                   WO200295027-A2
                                                                                                                                                                                                                                                                   modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                    Immunostimulant; oligodeoxynucleic acid; ODN; vaccine; DNA-RNA hybrid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New LS147 polypeptide,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5393 AAAAAATACAAAAAAGAAAAAA 5415
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ) BILLINGEL P.
COHENT T L.
COLPITTS T L.
COLPITTS T L.
PRIEDMAN P N.
KLASS M R.
RUSSELL J C.
STROUPE S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Stroupe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cohen M,
Stroupe S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97US-0048810P
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                                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                            *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     acid molecule ODN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
                                                                                                                                                                                                                     base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful for preparing a composition
                                                                                                                                                                                              "thiophosphate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 18.2; DB 1;
Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24.
                                                                                                                                                                                                   backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Friedman PN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1; Length
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RESULT 236
ACA62282/c
ID ACA622
XX ACA622
XX ACA622
XX Oligo
XX Oligo
XX Synthe
XX US2003
XX US2003
XX OJAN
XX US2003
XX US2003
XX WFI; 25-JAN
XX (EPIC-
XX WFI; 2
XX UMPI; 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         invention is based on the discovery that ODNs containing deoxyuridine residues (U-ODNs) have an immunostimulatory effect comparable to, or in many instances greater than, ODNs containing CpG motifs, producing higher numbers of specific T cells to a given antigen. The U-ODNs do not induce the systemic production of pro-inflammatory cytokines and, in contrast to CpG ODNs, are not dependent on a specific motif or a palindromic sequence. Use of a U-ODN for the preparation of a vaccine is claimed. Combining the U-ODN with an antigen strongly increases the potential of the antigen to raise the protection/immune response of a vaccinated individual. An example of the invention demonstrated the generation of a specific immune response against a melanoma-derived peptide (see ABP58360) by injection of mice with the peptide in combination with ODN
                                                                                                                                                                                 25-JAN-2000; 2000US-00494212
                                                                                                                                                                                                                07-SEP-2001; 2001US-00949305
                                                                                                                                                                                                                                                  30-JAN-2003
                                                                                                                                                                                                                                                                                US2003022318-A1
                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                               promoter
                                                                                                                                                                                                                                                                                                                                                                88; PCR;
                                                                                                                                                                                                                                                                                                                                                                                               Oligo (dT) primer #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                 12-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACA62282;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACA62282
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 26 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New oligodeoxynucleic acid molecules useful for the preparation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CIST-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence is that of a thiosubstituted oligodeoxynucleic acid (ODN) molecule, ODN 24, including deoxyuridine monophosphates. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2003-183880/18
                                                                                                                                                (EPIC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5393 AAAAAATACAAAAAGAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20;
                                                                                                                                                BPICLONE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ζ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INTERCELL BIOMEDIZINISCHE FORSCHUNGS. CISTEM BIOTECHNOLOGIES GMBH.
                                                                                                                                                                                                                                                                                                                                              primer; antisense therapy; \mathtt{mRNA} expression containing primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAAAAAAAAAAAAAAAAAA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 32;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.3%;
ilarity 87.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schellack C,
                                                                                                                                                                                                                                                                                                                                                                                                                                 (first
                                                                                                                                                  INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 0 C; 0 G; 1 T; 25 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Schmidt W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 18.2; DI
Pred. No. 3.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .7e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
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                                                                                                                                                                                                                                                                                                                                                                  profile;
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Improved polymerase thermocycling reaction for nucleic acid amplification, by thermal cycling of promoter-linked nucleic acid template synthesis and in vitro transcriptional amplification of racid sequences.

of nucleic

2003-479488/45

Ying

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Example 4; Page 14; 28pp; English

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RESULT 237
ADD44609/C
ID ADD4446
AC ADH446
AC ADH446
XX
DT 25-MAR
XX
DE Human
XX
KW Human;
KW Human;
KW Lumoux
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC (M1) for linear amplification of mucleic acid sequences, involved containing a number of mucleic acid sequences, involved containing a number of deoxynuclocide triphosphates and ribonucleotide triphosphates (1), combining the CC denatured (1) with a promoter-containing primer (P1), a primer (P2), a number of deoxynuclocide triphosphates and ribonucleotide triphosphates, contacting P2 with the denaturing the promoter-containing templates, contacting P2 with the denatured promoter-containing templates, contacting P2 with the denatured promoter-containing templates, contacting P2 with the denatured promoter-containing templates to generate a number of promoter-containing double-stranded nucleic acid templates are flanked by P1 in one contacting the promoter-containing double-stranded was templates to form a number of amplified RNA sequences, including the primer region of the promoter-containing double-stranded by P1 in one contacting the amplified RNA sequences with P2 to form a number of contacting the amplified RNA sequences with P2 to form a number of contacting the amplified RNA complified in a diseased state, and denaturing the bNA-RNA hybrid templates. The method is useful for improved polymerase thermocycling reaction for linear call successful for the promoter-containing double-stranded by M1 to generate multiple copies of the appreciation of mucleic acid sequences, and thus for producing mRNA composition profile by the calls in both normal and diseased states, comparing the expression profile of the cells in the normal and diseased states, comparing the call successful for treating the expression profile of the cells in the normal and diseased state, amplifying the expression profile of the cells in the normal and diseased state, amplifying the expression profile of the promoter protein, and the amplifying the antisense mRNA sequences by M1, and cell (s) in non-diseased state, amplifying the antisense mRNA sequences of a composition of the cell in the diseased state, amplifying the antisense mRNA sequ
                                                                                                                                                                                                                                                                                                              밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
Human; ss; Zalphail ligand; Zalphail receptor; immune response; tumour progression; metastasis; tumour stasis; haematopoietic tumour; lymphoma; B cell tumour; systemic lupus erythematosus;
                                                                                Human cDNA encoding Zalphall sequencing primer #3.
                                                                                                                                                                   ADH44609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 26 BP; 0 A; 0 C; 0 G; 26 T; 0 U; 0 Other;
                                                                                                                                                                                                          ADH44609 standard;
                                                                                                                                                                                                                                                                                                                                                      5393 AAAAAATACAAAAAGAAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                              AAAAAAAAAAAAAAAAAAAA 4
                                                                                                                           (first
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                                                                                                                                                                                                                                                                                                                                                                                                              0.3%;
                                                                                                                         entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          improved polymerase thermocycling reaction
                                                                                                                                                                                                                                                                                                                                                                                                              Score 18.2; DB 1;
Pred. No. 3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Length 26;
                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 238 ADI00945/c

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8
                                                                                                                                                                        Cc exposed to an antigen or pathogen comprising administering a composition comprising mature zalphall ligand polypeptide comprising residues 32-162 cc of ADH44572 in a pharmacutical vehicle. Also included are stimulating an immune response in a mammal exposed to an antigen or pathogen comprising zalphall ligand polypeptide in a pharmacutical vehicle; (c) comprising zalphall ligand polypeptide in a pharmacutical vehicle; (c) determining (in)directly the level of antigen or pathogen in the mammal; (d) administering a composition comparing the antigen or pathogen level in (a) with (b), where a change in the level indicates stimulation of immune response), and comparing the later indicates of immune response, and comparing (in) determining a newel of antigen or pathogen or pathogen or pathogen or pathogen (comprising: (a) determining a newel of antigen or pathogen or pathogen (comprising the level of the antibody; (d) comparing the level of the antibody in (a) with (b), where a cadministration level of the antibody in (a) with (b), where an comparing the level of the antibody in (a) with (b), where an comparing the level of the antibody in (a) with (b), where an comparing the level of the antibody in (a) with (b), where an comparing the level of the antibody in (a) with (b), where an comparing the level of the antibody in (a) with (b), where an comparing the level of the antibody in (a) with (b), where an comparing the level of the antibody in (a) with (b), where an comparing the level of the antibody in (a) with (b), where an comparing the level of the antibody in (a) with (b), where an activity resulting in a reduction in tumour progression, decrease in memmal the activity of the antibody in (a) with (b), where an activity of the antibody in (a) with (b), where an activity of the antibody in (a) with (b), where an activity of the antibody in (a) with (b), where an activity of the antibody in (a) with (b), where an activity of the antibody in (a) with (b), where an activity of the antibody in (a) with (b), where 
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                                                                            Matches
                                                                                                       Query Match
Best Local :
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11-MAR-1999; 99US-0123904P.
01-JUL-1999; 99US-0142013P.
09-MAR-2000; 2000US-00522217.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       rheumatoid arthritis; myasthenia gravis; diabetes; infectious immunocompromised patient; HIV infection; vaccine; primer.
                                                                                                                                                                      Sequence 26
                                                                                                                                                                                                              vaccines. The period exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to stimulating an immune response exposed to an antigen or pathogen comprises administering
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 7; SEQ ID NO 39; 103pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novak JE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-AUG-2001; 2001US-00923246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ZYMO ) ZYMOGENETICS
                                      5393 ААЛАЛАТАСАЛАЛАДАЛАЛАЛ 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                lating an immune response in a mammal exposed to an antigen or gen, useful for enhancing anti-tumor activity resulting in red progression or metastasis, comprises administering zalphall !
                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Johnston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Presnell
                                                                                                                                                                      BP; 0
                                                                                     Conservative
                                                                                                                                                                                                                of.
                                                                                                                                                                      A; 1 C; 0 G; 25 T; 0
                                                                                                                                                                                                                the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ź,ż
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INC.
                                                                                                       0.3%;
                                                                                                                                                                                                                   invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sprecher CA, Foster DC,
Nelson AJ, Dillon SR, I
                                                                                   0; Mismatches
                                                                                                       Score 18.2; DB 1
Pred. No. 3.7e+02
                                                                                                                                                                        U; 0 Other;
                                                                                                                           DB 1; Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hammond
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in a mammal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    zalphall ligand
                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in reduced
                                                                                     Gaps
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RESULT 239
ADO47862/c
ID ADO47862 standard; DNA; 26 BI
XX
AC ADO47862;
XX
DT 29-JUL-2004 (first entry)
XX
Gene expression inhibition as
XX
W gene expression; gene expreds
KW gene expression; viral pe
KW eukaryotic cell characterist;
XW microbial infection; viral pe
KW cancer cell proliferation; pc
XX
PN US2004087526-A1.
XX
PD 06-MAY-2004.
XX
PP 19-MAR-2003; 2003US-00393450.
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                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novak JE,
Gross JA,
                                                                                                            gene expression; gene expredssion innivition, eukaryotic cell characteristic; cell division rate; pigmentation; microbial infection; viral pathogenic infection; cancer cell proliferation; poly(dT)-26mer primer; ss; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel isolated zalphall ligand polypeptide. The polypeptide of the invention may be useful for boosting immunity to infectious diseases and treating immunoccompromised patients, such as HIV patients, as well as in improving vaccines. The current sequence is that of the PCR primer which was used in the exemplification of the invention.
                                                                                                                                                                                                   Gene expression inhibition associated poly(dT)-26mer primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New zalphall ligand polypeptides, useful for boosting immunity to infectious diseases, and treating immunocompromised patients, such as human immunodeficiency virus (HIV) patients, or in improving vaccines.
      19-MAR-2003; 2003US-00393450.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 26 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                09-MAR-2000; 2000US-00522217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  zalphall ligand; immunity; infectious disease; immunocompromised patient; HIV; vaccine; human; ss; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-NOV-2002; 2002US-00295723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-JUL-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequencing primer SEQ 39 used to analyse human zalphall ligand clone DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ZYMO ) ZYMOGENETICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADI00945 standard; DNA; 26
                                                                                                                                                                                                                                                                                                                                                                                                5393 АЛАЛАЛАТАСАЛАЛАСАЛАЛАЛ 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2003-811003/76
                                                                                                                                                                                                                                                                                                                                                                     25
                                                                                                                                                                                                                                                                                                                                                                                                                                 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                       AAAAAAAAAAAAAAAAAAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Presnell SR, Johnston JV,
                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ID NO 39;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; 1 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                              0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sprecher CA,
Nelson AJ, I
                                                                                                                                                                                                                                                                                             ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No. 3.76
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 18.2;
Pred. No. 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dillon SR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Foster DC,
                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hammond
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Holly
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                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                        cancer;
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The invention decribes a composition (I) for inhibiting the expression of a targeted gene in a substrate, comprising a DNA-RNA hybrid. (I) is useful for inhibiting the expression of the targeted gene in a substrate. The substrate is a prokaryote such as a viral or bacterial cell, or eukaryote or the cell of the eukaryote such as a vertebrate. The eukaryote is a mouse, rat, chimpanzee, preferably a human being. (I) is useful for altering the characteristics of an eukaryotic cell. The characteristic is chosen from expression of a protein, cell division rate and pigmentation. (I) has an effect that lasts at least three days. (I) is useful to inhibit the expression of messenger RNA is transcribed from a gene chosen from viral gene, oncogene, enzyme. (I) is useful for suppressing cancers, by knocking out cancer related genes, for preventing and treating microbial infections, preferably reducing viral pathogenic infection and for reducing the proliferation of cancer cells. This sequence represents a poly(dT)-26mer primer used in the creation of DNA-RNA hybrids for controlling gene
Query Match
Best Local S
Matches 20
                                                                           Sequence 26
                                                                                                                    expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 5; SEQ ID NO 6; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Composition useful for inhibiting the expression of a targeted gene in substrate, and for altering a characteristic of a eukaryote, comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-356242/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lin
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18-JAN-2002; 2002US-00052486.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA-RNA hybrid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (LINS/) LIN S.
(JIHH/) JI H H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ŝ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11 HH;
                    Similarity
                                                                             BP; 0
                                                                           A; 0 C; 0 G; 26
                  0.3%;
Score 18.2; DB 1
Pred. No. 3.7e+02
0; Mismatches
                                                                             T; 0
                                                                             U; 0 Other;
                                    DB 1; Length 26;
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D D

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RESULT 240
ADP19768/c
ID ADP197
XX ADP197
XX ADP197
XX Human
XX Cytost
KW Cytost
KW PCR; p
XX Homo s
XX US2004
PN US2004
XX US2004
PN US2004
PN US2004
XX IO-JUN
XX IO-MAI
PR 01-JUN
PR 01
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                                                    09-MAR-1999;
11-MAR-1999;
01-JUL-1999;
09-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cytostatic; zalphall ligand; pharmaceutical; cancer; immune response; melanoma; tumour; solid tumour; haematopoietic tumour; lymphoma; huma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human zalpha11 ligand
                                                                                                                                                                                                                                                                  10-SEP-2003; 2003US-00659684.
                                                                                                                                                                                                                                                                                                                                                                                                                                US2004110932-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADP19768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADP19768 standard;
                                                                                                                                                                                                                                                                                                                                                     10-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5393 AAAAAAATACAAAAAAGAAAAAA 5415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAAAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                       99US-0123904P.
99US-0142013P.
                                                                                                                                                                          99US-0123547P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR primer seqid 39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  26
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RESULT 241
ADQ80457
ID ADQ804
XX ADQ804
XX ADQ804
XX PA (26)
DT 09-SEP
XX RNA hy
KW Primer
XX Uniden
XX Uniden
XX Uniden
XX Uniden
XX Uniden
XX ON-DEC
XX OR-DEC
XX (GENO-PA (RUFF)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC a mammal; a method of stimulating an immune response in a mammal with CC melanoma; a method of stimulating an immune response in a mammal bearing CC a tumour; an isolated polynucleotide comprising a sequence of nucleotides CC that encode amino acid residues cited above, where the polynucleotide CC encodes a polypeptide that binds a receptor comprising 538 amino acids, CC fully defined in the specification; a pharmaceutical composition CC comprising the polynucleotide encoding, in a pharmaceutically acceptable CC vehicle; an expression vector comprising the following operably linked CC elements a control element; and a DNA segment comprising the CC polynucleotide; and an isolated polynucleotide molecule comprising at CC defined in the specification. The molecules, compositions and methods are CC useful for treating cancer, e.g. melanoma, solid tumour, haematopoietic CC tumour, or lymphoma. This sequence represents a primer used in the cx expression cloning of human cytokine zalphall ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novak JE,
Gross JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention describes an isolated polypeptide comprising a sequence of amino acid residues that is at least 90 or 95% identical to residues 41 (Gln) to 148 (Ile), or 32 (Gln) to 148 (Ile), or 32 (Gln) to 148 (Ile) of a sequence of 162 amino acids (SEQ ID NO:2, human zalpha11 ligand), fully defined in the specification. Also described are: a pharmaceutical composition comprising the polypeptide, and a vehicle; a method of treating cancer in
   Jimenez
                                                                                                                                                                                                                                           Unidentified
                                                                                                                                                                                                                                                                             primer
                                                                                                                                                                                                                                                                                              RNA hybrid; self normalisation; five prime exon rescue; RNA linker; ss;
                                                                                                                                                                                                                                                                                                                                 Da (26) biotin primer
                                                                                                                                                                                                                                                                                                                                                                       09-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                           ADQ80457;
                                                                                                                                                                                                                                                                                                                                                                                                                                               ADQ80457 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 26 BP; 0 A; 1 C; 0 G; 25 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New zalphall ligand polynucleotide and polypeptide molecules, useful treating cancer, e.g. melanoma, solid tumor, hematopoietic tumor, or
                                                                                             06-DEC-2002; 2002GB-00028557
                                                                                                                                                                   24-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 7; SEQ ID NO 39; 111pp; English
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                                                                                                                               08-DEC-2003; 2003WO-GB005341
                                                                                                                                                                                                     WO2004053160-A2
                                     (RUFF/)
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                                                       GENOMICA SAU
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Johnston JV,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                     (first entry)
 Escobar IG,
                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sprecher CA, Foster DC, Holly Nelson AJ, Dillon SR, Hammond
 Gallego SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 18.2;
Pred. No. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
 Cimadevilla JCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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RESULT 242
AL47121/c
ID AAL471
XX AAL471
XX AAL471
XX Pyrin
XX Pyrin
KW Pyrin
KW antiar
KW neurop
KW inflam
KW arteri
KW osteoa
XX Uniden
XX Uniden
XX Uniden
XX Uniden
XX WO2002
XX OS Uniden
XX WO2002
XX Uniden
XX WO2002
XX Uniden
XX WO2002
XX Uniden
XX Un
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Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to experimentally analysing boundaries within polymeric DNA or RNA molecules comprises hybridizing first and second different RNA molecules, derived from first and second samples, to provide RNA / RNA hybrids. The product of the method above is used as a probe. The method is useful in experimentally analysing boundaries within polymeric DNA or RNA molecules. The methods are useful in analysing polymeric nucleic acid sequence variations and in identifying molecules of therapeutic interest. The present sequence represents a RNA primer of the inventor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Experimentally analyzing boundaries within polymeric DNA or RN: molecules, useful in analyzing polymeric nucleic acid sequence variations, comprises hybridizing different RNA molecules to p://RNA hybrids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pyrin domain; PYD domain; antiinflammatory; antiparkinsonian; antiarteriosclerotic; antipsoriatic; antibacterial; virucide; neuroprotective; antiarthritic; antirhematic; antiasthmatic; nephrotropic; osteopathic; nootropic; intracellular signal transduction; inflammation; Alzheimer's disease; infection; psoriasis; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 26 BP; 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-OCT-2001; 2001WO-EP012545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unidentified.
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                                                           Example; Page 49; 116pp; German.
                                                                                                                                                    New DNA encoding protein with pyrin domain, useful for treating diseases involving impaired signal transduction, particularly inflammation, also
                                                                                                                                                                                                                                                                                                     Tschopp J, Martinon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pyrin domain containing protein coding sequence PCR primer JT1525.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the invention.
                                                                                                                                                                                                                                               WPI; 2002-427093/45.
                                                                                                                                                                                                                                                                                                                                                                    (APOT-) APOTECH RES & DEV LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                15-NOV-2000;
30-NOV-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-MAY-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200240668-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  osteoarthritis; glomerulonephritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 arteriosclerosis; multiple sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5393 ААААААТАСАЛАЛАДААААА 5415
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard; DNA; 18 BP
                                                                                                                           and antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAAAAAAAAAAAAAAAAAA 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                2000DE-01056687.
2000DE-01059595.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NO 14; 36pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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Pred. No. 3.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rheumatoid arthritis; sarcoidosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.7e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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RESULT 243
AAL47123
ID
ACC AAL471
XX AAL47
XX AA
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Best Local Similarity
                                                                                                              The present invention relates the DNA and their encoded proteins, where the proteins contain at least one PYD (pyrin) domain. These can be used to treat diseases associated with impaired intracellular signal transduction, particularly inflammation such as psoriasts, arteriosclerosis, bacterial or viral infections (particularly meningitis and pneumonia), multiple sclerosis, rheumatoid arthritis, asthma, sarcoidosis, glomerulonephritis and osteoarthritis, and also Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                        New DNA encoding protein with pyrin domain, useful for treating diseases involving impaired signal transduction, particularly inflammation, also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiarteriosclerotic; antipsoriatic; antibacterial; virucide; neuroprotective; antiarthritic; antitatheumatic; antiasthmatic; nephrotropic; osteopathic; nootropic; intracellular signal transduction; inflammation; Alzheimer's disease; infection; psoriasis; asthma; arteriosclerosis; multiple sclerosis; rheumatoid arthritis; sarcoidosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL47123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 18 BP; 5 A; 5 C; 5 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                proteins and
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                                                                                                                                                                                                                                                                                                                                                      Example; Page 49; 116pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (APOT-)
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                                                                                        and also Alzheimer's PCR primer used to
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Matches 18
                                                                                              The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding LIM domain kinase 1. The compound specifically hybridises with the nucleic acid molecule encoding LIM domain kinase 1 and inhibits the expression of LIM domain kinase 1. It specifically hybridises with at least an 8-nucleobase portion of a preferred target region on the nucleic acid molecule encoding LIM domain kinase 1. The antisense oligonucleotide is useful for modulating the expression of LIM domain kinase 1 in cells or tissues to treat diseases associated with their expression, such as a developmental disorder or a neurological disorder. In addition, the compound is used for diagnostics, prophylaxis, or as research reagents or kits. This sequence represents a treatment of the compound is sequence to the compound is the compound is compound to the compound is sequence to the compound is the compound is compound in the compound in the compound in the compound is compound in the compound in the compound in the compound is compound in the co
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotides for modulating expression, useful for diagnosing, preventing associated with the kinase, e.g. neurological
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 15;
                                                                           numan LIM domain kinase 1 antisense oligonucleotide.
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3.7e+02;
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Best Local Similarity
Matches 18; Conserv
                                            The invention describes a compound 8-80 nucleobases in length targeted to a nucleic acid molecule encoding LIM domain kinase 1. The compound specifically hybridises with the nucleic acid molecule encoding LIM domain kinase 1 and inhibits the expression of LIM domain kinase 1. It specifically hybridises with at least an 8-nucleobase portion of a preferred target region on the nucleic acid molecule encoding LIM domain kinase 1. The antisense oligonucleotide is useful for modulating the expression of LIM domain kinase 1 in cells or tissues to treat diseases associated with their expression, such as a developmental disorder or a neurological disorder. In addition, the compound is used for diagnostics, prophylaxis, or as research reagents or kits. This sequence represents a human time domain kinase 1 artisense of its or kits.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonucleotides for modulating LIM domain kinase expression, useful for diagnosing, preventing or treating condit associated with the kinase, e.g. neurological or developmental of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2004-121553/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-JUL-2002; 2002US-00199199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-JUL-2002; 2002US-00199199
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuroprotective; LIM domain kinase 1; developmental disorder; neurological disorder; diagnostic; prophylaxis; human; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human LIM domain kinase 1 antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADI38771;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2643 GCAGCTGCTGCTGCAGCC 2660
                             LIM domain kinase 1 antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
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                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 70; 81pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.3%; So
llarity 100.0%; I
Conservative 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "OTHER= 2'-O-Methoxyethyl (2'-MOE) nucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag= a
/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "OTHER= Phosphorothioate backbone.
are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2'-0-Methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Lo
3.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              or treating conditions or developmental disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>,</u>
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RESULT 246
ADJ13738/c
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                                                                                                       Query Match
Best Local S
Matches 18
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                           This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                     Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Analysis of chemical modification of DNA involves obtaining sample of
to be analyzed, treating DNA with chemical reagents that result in
different base sequences, and determining sequence of resulting DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Garner HR, Minna JD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-AUG-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; SEQ ID NO 865; 210pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tumour suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 probe; ss; chemical modification; methylation; array; CpG island
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (BALO/)
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                                                                                                                                       Local Similarity
                                                      2436 GGATGAGAAGGGGAGAGG 2453
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BALOG R P
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                                                                                                          Conservative
                                                                                                                                                                                                                     BP; 2 A;
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Conservative (
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                                                                                                                                                                                                                     11 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                 0.3%;
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                                                                                                          0; Mismatches
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Pred. No.
                                                                                                                                    Score 18;
Pred. No.
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                                                                                                                                                                 DB 1;
                                                                                                                                       3.8e+02;
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                                                                                                                                                                 Length 21
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                                                                                                             Indels
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                                                                                                          Gaps
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RESULT 248
AAT86582/c
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AAT85350/c
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                                                                                                                                                                                                                                                     encoding silk protein. The process involves: (a) selecting target DNA, from a silk-producing spider, that contains many repetitive and non-repetitive regions; (b) selecting a single-stranded DNA primer of at least 10 nucleotides with a sequence that is complementary to a region of the target; (c) repetitively combining the primer with melted target DNA, incubating the mixture with nucleotides and a DNA polymerase with containing the mixture with nucleotides and a DNA polymerase with the proofreading activity to produce a DNA fragment which is complementary to the target and is at least 2 kb long. The present sequence represents a specifically claimed primer for use in this process. The DNA fragment can be used to make fibres, films, woven articles, e.g. for use in parachutes, sails, body armour, and absorbers (e.g. of heavy metals, biological weapons, DNA, chemicals, flavours and fragrances). The high molecular weight (90-250 kD) of spider silk proteins can be produced on a commercial scale (at over 2 g/l cell mass). It has better tensile series and elasticity than silkworm silk. Inclusion of both repetitive
                                                                                                                                                                                  Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New opt. multimerised DNA sequences encoding spider silk protein - contg. both repetitive and non-repetitive sequences, useful for making high strength films, fibres, woven articles etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          High strength film; fibre; woven article; parachutes; sails; absorber; body armour; heavy metal; biological weapon; chemical; flavour; fragrance; Nephila clavipes; ss.
   AAT86582;
                                    AAT86582 standard; DNA; 21
                                                                                                                                                                                                                                                     Sequence 23 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 7; Page 54; 57pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Basel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A process has been developed for the production of a DNA fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22-AUG-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (BASE/) BASEL R M.
(BLIO/) ELION G R.
                                                                                                                                                   2640
                                                                                                                                                                                                                                                                                   non-repetitive regions ensures isolation of stable clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RM,
                                                                                                                     23
                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            target DNA primer
                                                                                                                                                                                                0.3%;
90.0%;
                                                                                                                                                                                                                                                  6 C; 10 G; 2 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23
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                                                                                                                                                                                    1;
                                                                                                                                                                                                 Score 18; DB 1;
Pred. No. 3.8e+02;
                                                                                                                                                                                    Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 249
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents a phosphorothioate oligonucleotide which was prepared by solid phase synthesis. The method comprises adding at least one dimeric phosphoramidite synthon, optionally having a protected thioseter group in its internucleotide link, during the synthesis cycle. These novel dimeric phosphoramidite synthons are used as antisense molecules for inhibition of gene expression. The method gives increased yields of the phosphorothiate oligonucleotide (since fewer cycles are needed) and facilitates separation of impurities (greater difference in size compared with use of monomeric synthons)
                            DNA fidelity; DNA manipulation; cancer; rragile a symmetry, compounds distribution of the station of the statio
                                                                                                             Endonuclease; MED1; human; methyl-CpG binding endonuclease-1; PCR primer;
DNA fidelity; DNA manipulation; cancer; fragile X syndrome; therapy;
                                                                                                                                                                                          PCR primer for human MED1 endonuclease coding sequence.
                                                                                                                                                                                                                                                    19-MAY-1999
                                                                                                                                                                                                                                                                                                    AAX22013;
                                                                                                                                                                                                                                                                                                                                                           AAX22013 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 0 A; 10 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; Page 20; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Solid phase synthesis of phosphoro:thioate oligo:nucleotide(s) using new dimeric synthon(s) - useful as anti:sense molecules for inhibiting gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1997-415290/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reese CB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-FEB-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Phosphorothicate
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Best Local &
                  Addison's disease; atrophic gastritis; degenerative nervous system disease; type I hypersensitivity; Alzheimer's disease; hypersensitivity disease; type I hypersensitivity; allergy; type II hypersensitivity; Goodpasture's syndrome; allergy; type II hypersensitivity; leprosy; infections disease; viral infection; type IV hypersensitivity; leprosy; infection; disease; viral infection; fundal infection; schaftosome; filaria; bacterial infection; Mycobacterium; neoplastic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  its prognosis, where caused by alterations in the MED1-encoding gene, are identified by sequence comparison, amplification, detecting altered polypeptide, and restriction fragment mapping, hybridisation (particularly to probes specific for a mutant allele). These same methods can also be used to diagnose fragile X syndrome and other diseases (e.g., myotonic dystrophy, Huntington's disease, spinocerebellar ataxia and Kennedy's disease) associated with triplet repeat expansion. The DNA, or its fragments, are used as probes and primers in the above diagnostic methods, also to isolate homologous sequences, as sources of antisense sequences and for gene transfer, particularly to restore drug sensitivity
                                                                                                                                                                         human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease;
                                                                                                                                                                                                                                                                                                                                                            ADH70613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP; 6 A; 3 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            maintaining DNA fidelity, for DNA manipulation and to raise antibodies. Susceptibility or predisposition to cancer (particularly colorectal or endometrial, especially hereditary non-polyposis colorectal cancer), or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This sequence is a PCR primer for DNA encoding the human MED1 endonuclease of the invention. MED1 (methyl-Cpg binding endonuclease-1) is used to screen for specific modulators (potential therapeutic agents particularly mimetics of MED1) and to study interactions involved in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 45; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid encoding human endonuclease MED1 involved in DNA mismatch repair - used for diagnosing susceptibility to cancer and fragile X syndrome, and therapeutically.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1999-142462/12.
                                                                                                                                                                                                                                                                   Human Vbeta gene repeat sequence #403.
                                                                                                                                                                                                                                                                                                                                                                                                     ADH70613 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      drug-resistant cancer cells
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Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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    cancer
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RESULT 251
AAQ34355
ID AAQ343
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XZ AAQ343
XX PCR; 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a kit for diagnosing and treating T-cell CC associated diseases which comprises a panel of nucleic acid primers composited by priming and allowing amplification of each Vbeta gene, CC VbetaRNA or cDNA. The kit is useful for diagnosing organ transplant Crejection and diagnosing and treating T-cell associated diseases. CC including autoimmune diseases, degenerative nervous system diseases, cand neoplastic diseases. Autoimmune diseases include Addison's diseases. CC and neoplastic diseases. Autoimmune diseases include Addison's diseases. CC atrophic gastritis. Degenerative nervous system diseases include Type CC atrophic gastritis. Degenerative nervous system diseases include Type CC sclerosis and Alzheimer's disease. Hypersensitivity diseases include Type CC Inpersensitivities such as contact with allergens that lead to CC allergies, Type II hypersensitivities such as those present in CC Goodpasture's syndrome and Type IV hypersensitivities such as those caused by the yeast genus Candida, parasitic infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by CC Schistosomes, filaria and bacterial infections such as those caused by CC Schistosomes, filaria and bacterial infections such as those caused by CC Schistosomes, filaria and bacterial infections such as those caused by CC Schistosomes, lymphomas and cancers such as cancer of the brain, CC breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                   Matches 19;
                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 0 A; 10 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                             AAQ34355 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comprises nucleic acid primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hood LE,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Vbeta gene
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(ROWE/)
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19-SEP-1995;
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95US-00531241.
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                                                                                                                                                                                                             DNA;
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Pred. No. 4e
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                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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genetic mapping; traits; amplification; ss

Upstream PCR primer TGLA351UP1

PCR; selection; microsatellite; OPTIPRIM; breeding; cattle; parentage;

25-MAR-2003 02-FEB-1993 AAQ34355;

(revised) (first en

entry)

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RESULT 252
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AC AAT764
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       microsatellites is summarised in the specification and indexed herein (see below). For each such microsatellite sequence sufficient information was obtd. to generate the required PCR primers for in vitro amplification of the corresp, microsatellite (using the program OPTPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding. See also AAQ33501-34440. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                         16-SEP-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 22 BP; 5 A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The sequence shows an upstream PCR primer for in vitro amplification of bovine microsatellite sequences obtd. by screening library of bovine MoJ DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 clones cross- hybridised. Assuming oligonucleotide probe one out of 50 clones and MboJ sites, the frequency of (T6)n > 9 microsatellites in the bovine genome is estimated at >100,000. The sequence information for ca. 230 such bovine
                                                                                                                                                            Synthetic.
                                                                                                                                                                                        Asthma; airway epithelium; adenosine free; cystic fibrosis; chronic obstructive pulmonary disease; bronchitis; ss.
                                                                                                                                                                                                                                        Chymase antisense oligonucleotide
                                                                                                                                                                                                                                                                                                        AAT76462;
                                                                                                                                                                                                                                                                                                                                      AAT76462 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Polymorphic bovine DNA markers - used in genetic identification, gene mapping, and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Georges M,
                                07-JUN-1995;
                                                                06-JUN-1996;
                                                                                               19-DEC-1996.
                                                                                                                           WO9640162-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Table
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9213102-A1
   (UYEC-) UNIV EAST CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 5069 CTCATCTGGTGGCCACAGCAG 5089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8; Page 471; 517pp;
                                                                                                                                                                                                                                                                                                                                                                                                                     N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                   CACATCTGGTGGCCACATCAG 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             маввеу JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.3%;
ilarity 90.5%;
Conservative
                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            9108-00642342
                                95US-00474497
                                                                96WO-US009306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92WO-US000340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 17.8;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.1e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 253
AAXS4254/c
ID AAX542
XX AAX542
XX AAX542
XX AAX542
XX AATise
AC Chymas
XX Antise
KW Impair
KW pulmon
KW respir
KW pulmon
KW pulmon
KW pulmon
KW pulmon
KW pulmon
KW pulmon
KW posta
XX Synthe
XX PO9913
XX Synthe
XX IT-SEP
PR 17-SEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      which involves the topical administration of an essentially adenosine free antisense oligonucleotide (ON) to the airway epithelium of the subject. The present sequence is an antisense oligonucleotide specific for chymase. The method can be used to treat airway diseases such as cystic fibrosis, asthma, chronic obstructive pulmonary disease, bronchitis and other airway diseases characterised by an inflammatory response. By eliminating adenosine from the antisense ON, its liberation upon antisense degradation is prevented, thereby preventing adenosine-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; altergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasoconstriction; emphysema;
  New antisense oligonucleotides used vasoconstriction.
                                                                                 WPI; 1999-229400/19.
                                                                                                                                                                                                                                            17-SEP-1997;
09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          colon cancer; breast cancer; lung hepatocellular carcinoma; kidney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chymase antisense oligonucleotide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX54254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX54254 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 22 BP; 0 A; 10 C; 2 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A method for treating airway disease in a subject has been produced, which involves the topical administration of an essentially adenosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 5; Page 41; 71pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Treatment of airway diseases such as asthma - by topically applying adenosine-free antisense oligo:nucleotide to airway epithelium of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1997-051871/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Мусе Ј₩,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1
                                                                                                                                                                                                                                                                                                                              17-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9913886-A1
                                                                                                                                                                                         (UYEC-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      148 CAGGACCCAGGAGAGGGAAGA 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 Similarity
19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bronchoconstriction in patients with hyper-reactive airways
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAGGAGACAGGAGAGGGAAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Metzger WJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                            EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                               97US-0059160P.
98US-00093972.
                                                                                                                                                                                                                                                                                                                              98WO-US019419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 17.8; DB 1; Pred. No. 4.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer; melanoma; hepatic metastasis;
                              ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2:
                              ę,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22
                              pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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treatment

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RESULT 254
ANA33698/c
ID ANA336
XX ANA336
XX Low ad
XX Low ad
XX Human;
KW phosph
KW allerg
KW antiall
KW pulmon
KW pulmon
KW cancex
XX Homo s
XX Wollon
PN WO2000
XX O3-AUC
XX O3-AUC
XX New al
PT Vasocc
PT Vasocc
PT Vasocc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC directed against at least 2 mRNAs selected from target genes, coding and CC conon-coding regions of RNAs corresponding to target genes, gene initiation CC codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3' CC -end and the juxta-section between coding and non-coding regions and all CC segments of RNAs encoding proteins associated with one or more diseases, CC conditions or mixtures. The antisense oligonucleotides may be derived CC from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-271) can be used for the antisense treatment of CC diseases and conditions. Typical diseases and conditions are those CC diseases, pulmonary vasoconstriction, inflammation, including lung CC diseases, pulmonary vasoconstriction, inflammation, respiratory CC distress syndrome, pain, cystic fibrosis, pulmonary hypertension, cc pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary CC disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g. Cc colon cancer, breast cancer, lung cancer, melanoma, hepatic metastases, as compared to the lungs of cancers which may metastasize or have metastasized concerts and concerts and concerts of the concerts of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 19
New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothioate; impaired respiration; inflammation; allergy; disease; bronchoconstriction; inhibitor; antinflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cytotic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pulmonary hypertension; chronic cancer; leukaemia; lymphoma; can
                                                                                                              WPI; 2000-205971/18.
                                                                                                                                                                                                                                                                            03-AUG-1998;
                                                                                                                                                                                                                                                                                                                              03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200009525-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Low adenosine antisense oligonucleotide SEQ ID NO:1387.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA33698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA33698 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 0 A; 10 C; 2 G; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to the lungs,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 60; 120pp; English
                                                                                                                                                                                                                        (UYEC-) UNIV EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The specification describes antisense oligonucleotides (AAX52869-X55271)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   148 CAGGACCCAGGAGAGGGAAGA 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CAGGAGACAGGAGAGGGAAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   including breast and prostate cancer
                                                                                                                                                                                                                                                                            98US-0095212P
                                                                                                                                                                                                                                                                                                                              99WO-US017712.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lymphoma; carcinoma; metastasis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 17.8; DB 1;
Pred. No. 4.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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AAF19820/c
ID AAF198
XX Low ad
KW Low ad
KW human;
KW surfac
KW respir
KW surfac
KW respir
KW cancer
XX Cancer
XX Cancer
XX Homo s
XX Homo s
XX PPD 26-OCT
XX 24-MAR
XX 06-APR
XX 07EC-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cc e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, cimpeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, emphysema, chronic obstructive cc fibrosis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, carcinomas, and cancers which may metastasise to the lungs, including threast and prostate cancer. The reduction of the adenosine content of the CC ons reduces side effects. The A-containing ONs break down with the release of deoxyadenosine which activates adenosine receptors causing throughout the receptors causing throughout the present the control of the present the nucleotide sequences given in the sequence listing from the present control of the previously named sequences. SEQ ID NO:1 to 2815, and then the last 185 capuances are also called SEQ ID NO:1 to 185, but the sequences differ from the previously named sequences. SEQ ID NO:11 to 1680 (AAA32323 to CAAA3392) are specifically claimed ONs from the present invention on the match up with their corresponding SEQ ID NO: sequences given in the sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a new composition comprising an antisense oligonucleotide (ON) with low adenosine (up to 15%), which targets nucleic acids involved in bronchoconstriction, allergies, and/or inflammation. The ON can have antiinflammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compositions are useful for the treatment of diseases associated with inflammation, impaired aliways, including lung disease and diseases whose secondary effects afflict the lungs of a subject. They can be used for treating
                                                                                                                                                                                                                                                                                                                    human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection;
                                               06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human chymase polynucleotide fragment #1387.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF19820 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancers.
                                                                                            24-MAR-2000; 2000WO-US008020.
                                                                                                                                            26-OCT-2000
                                                                                                                                                                                      WO200062736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Low adenosine
                                                                                                                                                                                                                                 sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  148 CAGGACCCAGGAGAGGGAAGA 168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 CAGGAGACAGGAGAGGGAAGA 1
                                                                                                                                                                                                                                                                              88.
                                                                                                                                                                                                                                                                                                    obstructive pulmonary disease; pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 438; 1343pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antisense oligonucleotide; phosphorothioate; allergy;
                                                 99US-0127958P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10 C; 2 G; 10 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 17.8; DB 1; Length Pred. No. 4.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22;
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(UYEC-) UNIV EAST CAROLINA

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RESULT 256
AAF99703/c
ID AAF997
XX AAF997
XX AAF997
XX Immuno
XX Vaccin
KW Immuno
KW Immuno
KW Immuno
KW Infect
XX Yorthe
XX WO2001
XX WO2001
XX WO2001
XX WO2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C(I) can have respiratory, bronchodilator, antiinflammatory, analgesic, cimmunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC Immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. CC immunosuppression and or activity of target polypeptides associated with CC lung/respiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, immunoglobulins and antibodies, antibody receptors, cytokines and CC chemokines, endogenously produced specific and non-specific enzymes, binding proteins, adensine receptors, bandykinin receptors, central nervous system (CNS) and peripheral nervous and non-nervous system CC receptors, defensins, growth factors, vasociated proteins. The CC antidense oligonuclectides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary peptides and construction sufferent hypoproduction which are associated with a disease or cc condition selected from pulmonary vasoconstriction, inflammation, allergy(ies) and/or cancer. AAP18434 to AAP21543 represent human polynuclectide fragments and antisense oligonuclectides used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                 Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base (I) can have respiratory, bronchodilator, antiinflammatory, analgesic,
                                                                                                                                                                                                                                                          12-JUN-2001
                                                                                                                                                                                                                                                                                                   AAF99703
                                                                                                                                                                                                                                                                                                                                        AAF99703 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 0 A; 10 C; 2 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 14; Page 250; 1592pp; English.
                                         WO200122972-A2
                                                                                                                                                                                                                  Immunostimulatory nucleic acid #819.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes low adenosine (A) content antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       adenosine receptors during metabolism, useful e.g. for treating cancers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Low adenosine (A) content antisense oligonucleotides which do not trigger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NYCE/) NYCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    respiratory obstructions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              148 CAGGACCCAGGAGAGGGAAGA 168
                                                                                                                                                                                                                                                                                                                                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                                                                                                                                                          ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.8; DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 4.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
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ABSTULT 257
ABS78424/c
ID ABS784
AC ABS784
AC ABS784
AC ABS784
AC ABG109
T 13-DEC
XX Angiog
KW Angiog
KW tumour
KW diabet
KW cornea
KW rubeas
KW plaque
KW angiof
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX I1-JUL
XX 14-DEC
XX 14-DEC
XX 14-DEC
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothioate backbone
                                                                                                                                                                                  diabetic retinopathy, retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-1999; 99US-0156113P
27-SEP-1999; 99US-0156135P
23-AUG-2000; 2000US-0227436P
                                      14-DEC-2001; 2001WO-US048458
                                                                                                                                                                                                                                                                                               Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis;
                                                                                                                                                                                                                                                                                                                                                         Angiogenesis inhibitory oligonucleotide #908.
                                                                                                                                                                                                                                                                                                                                                                                              13-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                   ABS78424;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABS78424 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 22 BP; 0 A; 11 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 101; Page 56; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-273485/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Krieg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-SEP-2000;
                                                                                                                WO200253141-A2
                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for stimulating an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (IOWA ) UNIV IOWA RES FOUND (COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1180 AGAGAAAGAGAGAGAGAAA 1200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AGAGAGAGAGAGAGAGA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Schetter C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000WO-US026383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprises administering an immunostimulatory nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vollmer J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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14-DEC-2000; 2000US-0255534P

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RESULT 258
ACH03242/c
ID ACH032
XX ACH032
XX Inmunc
XX Inmunc
XX Inmunc
XX Inmunc
XX Inflan
XX US200:
XX US200:
XX US200:
XX US200:
XX US200:
XX IS-MAI
XX I
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, it more metastasis, precancerous lesion, rheumatoid arthritis, psoriasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunostimulatory; antiinflammatory; dermatclogical; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound gramulation, intestinal adhesions, atherosclerosis, scleroderma a hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
   disease by
                            Treating non-allergic inflammatory diseases, such as psoriasis, ecz allergic contact dermatitis, latex dermatitis or inflammatory bowel
                                                                                                                                                                                                                                                                                                                                                                                                            29-MAR-2002; 2002US-00112653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2003050268-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACH03242 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22
                                                                                                                                                                                                                                                                                                                                               29-MAR-2001; 2001US-0279642P
                                                                                                                            WPI; 2003-521815/49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunostimulatory nucleic acid #877.
                                                                                                                                                                                                                                                                                   (KRIE/) KRIEG A M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (COLE-) COLEY PHARM GROUP INC.
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                                                                                                                                                                                                                                                      BERG D J.
contact dermatitis, by administering an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGAGAGAGAGAGAGAGAGA 2
                                                                                                                                                                                      Berg DJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
   immunostimulatory
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11 T; 0
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No. 4.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U; 0 Other;
      nucleic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22;
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RESULT 259
ADB37205/c
ID ADB372
XX ADB372
XX ADB372
XX ADB372
XX Immunc
XX ds; al
KW ds; al
KW hypo-r
XX Synthe
XX US2003
XX US2003
XX O2-FEE
XX YA (FCUR)
PA (FCUR)
PA (FCUR)
PA (FOUR)
XX WPI; 7
XX Treat:
PT Treat:
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Query Match
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Matches 19
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Best Local S
Matches 19
                                                                                                                                                      The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents an immunostimulatory nucleic acid of the invention.
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                                                                                                         Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                      Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
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                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 17; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-657977/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bratzler RL,
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(PETE/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunostimulatory nucleic acid #819
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADB37205 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    allergy; asthma; poly-G nucleic acid; aerosol formulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1180 AGAGAAAGAGAGAGAGAAA 1200
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PETERSEN
FOURON Y.
                              Similarity
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                                                                                                       BP; 0 A; 11 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 0 A; 11 C; 0 G; 11 T; 0 U; 0 Other;
  Conservative
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                         0.3%;
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                           Score 17.8;
Pred. No. 4.
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Pred. No. 4.1e+02;
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     Mismatches
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                           4.1e+02
                                                       DB 1;
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                                                       Length
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1180 AGAGAAAGAGAGAGAGAAA 1200

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RESULT 260
ADC13691/c
ID ADC136
XX ADC137
XX ADC137
XX ADC14
XX ADC14
XX ADC14
XX ADC14
XX ADC16
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11-JUL-2001; 2001US-0304502P.
12-JUL-2001; 2001US-0305501P.
13-JUL-2001; 2001US-0305262P.
17-JUL-2001; 2001US-0305262P.
24-JUL-2001; 2001US-030635P.
27-JUL-2001; 2001US-030828P.
27-JUL-2001; 2001US-030827P.
01-AUG-2001; 2001US-0309255P.
10-AUG-2001; 2001US-0311753P.
19-SEP-2001; 2001US-0311753P.
25-FEB-2002; 2002US-0358932P.
05-MAR-2002; 2002US-0358932P.
05-MAR-2002; 2002US-0361765P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NOVX; FADD interacting protein; ATPase; H+ Transporting; Lysosomal; FGF 17; Single Pass Transmembrane; Beta-Ketcacyl Synthase; Neuralin 2; Glutamate Receptor Interacting Protein 2; ChR-Methyltransferase; NP25 Variant; GTPase-Activating Protein; ELKS; Sim2; RhogAP; Phospholipase; Scavenger Receptor Domain Containing Protein; Metallothionein IA; NOGO receptor; FYVE; NOELIN; Cyclin Regulatory Subunit; Tetratrico Peptide Repeat Protein; Cyclin Regulatory Subunit; Tetratrico Peptide Repeat Containing Protein; Phospholipanine; Histidine Ammonia-Lyase; Cellular Retinaldehyde-Binding; Phenylalanine; Histidine Ammonia-Lyase; Cellular Retinaldehyde-Binding; Glutamine Repeat Containing Protein; TWP Receptor Associated Factor2; Vacuolar Protein Sorting Homologue R-VPS3AA; Bola Domain Containing Protein; TWP Receptor Associated Factor2;
   The invention relates to a novel isolated NOVX polypeptide comprising: a sequence of 57-1149 amino acids as defined in the specification, or its mature form; a sequence that is at least 95% identical to the 57-1149 amino acid polypeptide; or a sequence comprising one or more conservative substitutions in the 57-1149 amino acid polypeptide. The NOVX proteins of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Guo X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-JUL-2002; 2002WO-US021359
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RAL Guanine Nucleotide Dissociation Stimulator; Armadillo/Beta-Catenin
Metalloprotease; TlO Ser/Thr-rich; Ring finger-like; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human NOVX probe,
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                                                                                                                                                                                                                                             preventing
                                                                                                                                                                                                                                                                                New NOVX polypeptide, useful for preparing a composition for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                     Malyankar UM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patturajan M,
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                                                                                                                                                                                    37; Page 303; 393pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Casman
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                                                                                                                                                                                                                                                                                                                                                                                                                            Gerlach VL, Ander
SJ, Hjalt T, Mili
Zhong M, Padigaru
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[, Miller
sdigaru M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                  CB,
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Kekuda R, Shimkets
L, Shenoy SG, Gorma
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RESULT 261
ABZ95514/c
ID ABZ955
XX ABZ955
XX ABZ955
XX IT-OCT
XX Human;
KW Human;
KW antiae
KW antiae
KW adenoe
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Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   preventing cancer. This polynucleotide sequence gene encoding a NOVX protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200285308-A2.
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mes 19; Conserv
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Pred. No. 4.1e+02
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24-APR-2001; 2001US-0286137P 23-APR-2002; 2002WO-US013135 (EPIG-) 2003-229219/22. ß **BPIGENESIS PHARM INC** Li Y, Tang ŗ Sandrasagra A, L, Shahabuddin ŝ Katz E, Pabalan Ç Aguilar

Zerhusen BD; himkets RA;

õ

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding kNAs, and glucocorticoid or non-glucocorticoid steroid or Disclosure; SEQ ID NO 10756; 872pp; English ubiquinone.

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RESULT 262
ABD19659/c
ID ABD196
XX ABD196
XX ABD196
XX ABD196
XX UAMAAN;
KW PESPIX
KW PESPIX
KW PULMON
XX PULMON
XX HOMO B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Loc
Matches
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                                                                                                                                                                                                                                                                          Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; allergic rhinitis; pulmonary hypotrension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
Pharmaceutical composition for treating asthma, has antisense oligonuclectide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.
                                                                                                                                                                                                    WPI; 2003-093058/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-APR-2001; 2001US-0286036P
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                                                                                                                                                                                                                                                                          Li Y,
Tang
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                                                                                                                                                                                                                                                                              Sandrasagra A,
5, Shahabuddin
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Pred. No. 4.1e+02;
                                                                                                                                                                                                                                                                              Katz E,
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RESULT 263
ADK61705/c
ID ADK617
XX ADK617
XX ADK617
XX DE Base C
XX YICE V
XX YICE V
XX JP2003
XX JP2003
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XX JP2003
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XX HOKU-
PR 02-MAY
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Base containing SSR sequence #9
                                                                   WPI; 2004-003560/01
                                                                                                                                                                                                                                                                                                                                                                                                                                   JP2003319782-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allerg reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 15;
                                                                                                                                                                                                                                                                                                  02-MAY-2002; 2002JP-00130645.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rice variety; amplification genetic marker; ds.
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                                                                                                                                   (HOKU-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          148
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                                                                                                                                   HOKKAIDO GREEN
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Pred. No. 4.
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Identifying rice variety using

base sequence containing SSR sequence

amplifying genetic marker.

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ARESULT 264
ADX617
XX
ADX617
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AC ADX617
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DT 06-MAY
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DT 06-MAY
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                                                                                                                 Matches
                                                                                                                                         Query Match
Best Local &
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Best Local
                                                                                                                                                                                                                                                                             The present invention relates to identifying a rice variety as amplification genetic marker and identifying whether test rice variety is any one of the 32 rice varieties e.g., Kasalath, breath which came or Hayamasari, Italica Livorno, Dunghan Shali, Arroz Da Terra, Fany, USSR22, Nihonbare. The method is useful for identifying rice variety and identifies excellent rice variety. The present sequence represents a base containing SSR sequence of the invention.
                                                                                                                                                                                                                             Sequence 22 BP; 11 A; 0 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 65; SEQ ID NO 17; 30pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2003319782-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oryza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        rice variety; amplification genetic marker; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Base containing SSR sequence #17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADK61713;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADK61713 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to identifying a rice variety as amplification genetic marker and identifying whether test rice variety is any one of the 32 rice varieties e.g., Kasalath, breath which came or Hayamassari, Italica Livorno, Dunghan Shali, Arroz Da Terra, Fany, USSR22, Nihombare. The method is useful for identifying rice variety and identifies excellent rice variety. The present sequence represents a base containing SSR sequence of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-MAY-2002; 2002JP-00130645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-MAY-2002; 2002JP-00130645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22 BP; 0 A; 11 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 34; SEQ ID NO 9; 30pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Identifying rice variety using base sequence containing SSR sequence amplifying genetic marker.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-003560/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HOKU-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1180 AGAGAAAGAGAGAGAGAAA 1200
                                                     1180 AGAGAAAGAGAGAGAGAAA 1200
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                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HOKKAIDO GREEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HOKUREN NOGYO KYODO KUMIAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGAGAGAGAGAGAGAGAGA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                         90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BIO KENKYUSHO KK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                           0
                                                                                                                                                                    Score 17.8;
                                                                                                              Mismatches
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                                                                                                                                            4.1e+02
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4.1e+02;
                                                                                                                                                                    DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                              Indels
                                                                                                           0,
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                                                                                                           Gaps
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RESULT 266
AAX00525/c
ID AAX005
XX
AC AAX005
XX
DT 30-MAR
XX
DT 30-MAR
XX
DE Antise
                                                                                                                                                                                                                                                                                                                                    The sequence is that of a bovine microsatellite sequence obtd. by careening a genomic library of bovine MboI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (T6)n >9 microsatellites in the bovine genome is estimated at >100, 000. The sequence information for ca. 230 such bovine microsatellites is summarised in the specification and indexed herein (see below). The sequences upstream and downstream of the microsatellite sequence were used to generate the required PCR primers for in vitro amplification of the corresp.

C microsatellite (using the program OPTIPRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                               Query Match
Best Local 8
                                                                                                                                                                                                                                                                                                                             breeding.
field.)
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                                                              AAX00525;
                                                                                             AAX00525
                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Table 7; Page 132; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mapping,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic bovine DNA markers - used in genetic identification, gene mapping, and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Georges M, Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genetic mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR; selection; primers; OPTIPRIM; breeding; cattle; parentage; genetic mapping; traits; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence of microsatellite from clone AGLA209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ33511;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JAN-1992;
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                                                                                                                                                                                                   1180 AGAGAAAGAGAGAGAGAAA 1200
                                                                                                                                                                      Н
                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                             23
                                                                                             standard; DNA;
                                                                                                                                                                      AGAGAGAGAGAGAGAGAGA 21
                                                                                                                                                                                                                                                                                             BP; 12 A; 0 C; 11 G; 0
                                                                                                                                                                                                                                   Conservative
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(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91US-00642342
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                                                                                                                                                                                                                                                0.3%; Score 17.8; DB 1; Length 23; 90.5%; Pred. No. 4.1e+02;
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                                                                                             24
                                                                                             ВP
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                                                                                                                                                                                                                                                                                             T; 0 U; 0 Other;
                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                   0
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0

Antisense oligonucleotide for poly-purine target sequence.

30-MAR-1999

(first entry)

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RESULT 267
AAXOO527
ID AAXOO5
XX AAXOO5
XX AOOO5
XX 30-MAR
DT 30-MAR
XX Antise
XX Intera
XX intera
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX OS-JA1
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This oligonucleotide represents an antisense oligonucleotides (ASO) targeted to a poly-purine mRNA sequence generated by a method of selectively ranking nucleic acid molecules for inhibitory efficiency. The method comprises: (a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence zone RNA:ASO-DNA hybrid nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid base pair fractions into formulas to determine the fractions of each of a series of 13 nearest-neighbour nucleic acid base pair types to provide determined fractions; and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types by a stability ranking to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligomer binding interactions to select sequence zones for antisense targeting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Selectively ranking nucleic acid molecules, for inhibitory efficiency comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 24 BP; 0 A; 12 C; 0 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-OCT-1994;
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                                                     US5856103-A.
                                                                                       Synthetic.
                                                                                                                       Target; antisense; selective rank; inhibition; ranking; stability; interaction; ss.
                                                                                                                                                                         Antisense oligonucleotide for poly-pyrimidine target sequence
                                                                                                                                                                                                             30-MAR-1999
                                                                                                                                                                                                                                                                               AAX00527 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Col 13-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1999-105098/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-MAR-1997;
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interaction; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (TEXA ) UNIV TEXAS
                                                                                                                                                                                                                                                                                                                                                                                                   1180 AGAGAAAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                                                                                   19;
                                                                                                                                                                                                                                                                                                                                                                   24
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                           (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94US-00320507
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  72pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 17.8;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     .1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Length 24;

0

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targeted to a poly-pyrimidine mRNA sequence generated by a method of selectively ranking nucleic acid molecules for inhibitory efficiency. The method comprises: (a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence zone RNA:ASO-DNA hybrid nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid base pair fractions into formulas to determine the fractions of each of a series of 13 nearest-neighbour nucleic acid base pair types to provide determined fractions; and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types by a stability ranking to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligomer binding interactions to select sequence zones for antisense targeting
                                                                                                                                                                                                                                                                                                                                                                                                                                        Selectively ranking nucleic acid molecules, for inhibitory efficiency comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions
  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clark CL,
                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Col 13-14; 72pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-MAR-1997;
                                                                                                                                                                                                                                                                                                                              This oligonucleotide represents an antisense oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (TEXA ) UNIV TEXAS.
24 BP; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gray DM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94US-00320507.
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  A; 0 C; 12 G; 0 T; 0 U;
  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                               and
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AAXOOS24

ID AAX

XX AAC AAAX

AX Tai

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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Target sequence #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX00524 standard; mRNA; 24
                                                                                                                                                                                                                                                                                                                                                                   03-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Target; antisense; selective rank; inhibition; ranking; stability;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    interaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1180 AGAGAAAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGAGAGAGAGAGAGAGA 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                       97US-00808474.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for antisense oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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Pred. No. 4.
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WPI; 1999-105098/09

Clark CL,

Gray DM;

(TEXA) UNIV TEXAS.

94US-00320507

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Selectively ranking nucleic acid molecules, for inhibitory efficiency - comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions and multiplying.
Disclosure, Col 13-14; 72pp; English
```

This sequence represents a target mRNA for the generation of antisense oligonucleotides (ASO) in a method of selectively ranking nucleic acid molecules for inhibitory efficiency. The method comprises: (a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence zone RNA:ASO-DNA hybrid nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid sequence; (c) substituting nearest-neighbour nucleic acid base pair fractions into formulas to determine the fractions of each of a series of 13 nearest-neighbour nucleic acid base pair types to provide determined fractions; and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types by a stability ranking to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligomer binding interactions to select sequence zones

Sequence 24 BP; 12 A; 0 C; 12 G; 0 T; 0 U; 0 Other; 멂

antisense targeting

밁 ঠ Query Match Best Local S Matches 19 Local Similarity 1180 AGAGAAAGAGAGAGAGAGAAA 1200 AGAGAGAGAGAGAGAGAGA 21 Conservative 90.5%; 0 Score Pred. Mismatches No. 17.8; .1e+02 1; Length 24; Indels 0 Gaps

AAX00526 standard;

mRNA;

Poly-pyrimidine target sequence for antisense oligonucleotides 30-MAR-1999 (first entry)

Target; antisense; selective rank; inhibition; ranking; stability; interaction; 88.

US5856103-A

05-JAN-1999

03-MAR-1997; 97US-00808474

94US-00320507

(TEXA) UNIV TEXAS

Clark CL, Gray DM;

RESULT 269
AXXO0526/c
ID AXXO05
XX AXXO05
XX AXXO05
XX JO-MAR
XX Target
KW Intera
XX Synthe
XX Synthe
XX O5-JAN
XX O5-JAN
XX O7-OCI
XX O7-OCI
XX O7-OCI
XX OTEXA
XX O Selectively ranking nucleic acid molecules, for inhibitory efficiency comprises determining the fraction a set of nearest-neighbour nucleic acid base pair types in a target sequence zone, substituting nearest-neighbour nucleic acid base pair fractions to determine the fractions

Disclosure; Col 13-14; 72pp; English

This sequence represents a target mRNA for the generation oligonucleotides (ASO) in a method of selectively ranking of antisense nucleic acid

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molecules for inhibitory efficiency. The method comprises: (a) determining the fraction of each of a set of 13 nearest-neighbour nucleic acid base pair types in a target sequence zone RNA:ASO-DNA hybrid nucleic acid sequence; (b) substituting nearest-neighbour nucleic acid base pair fractions into formulas to determine the fractions of each of a series of 13 nearest-neighbour nucleic acid base pair types to provide determined fractions, and (c) multiplying the fractions of the 13 nearest-neighbour nucleic acid base pair types by a stability ranking to the nucleic acid antisense sequence; where the results are ordered to produce a ranking. The process is used to rank nucleic acid sequences based on the stability of nucleic acid oligomer binding interactions to select sequence zones for antisense targeting
Sequence 24
BP; 0 A; 12 C; 0
G; 0 T; 12 U; 0 Other;
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Query Match Best Local S Matches 19 Local Similarity 19; Conservative 90.5%; 0 Score 17.8; DB 1 Pred. No. 4.1e+02 Mismatches ۲. Length Indels 24 0 Gaps 0,

문 Ś 1180 AGAGAAAGAGAGAGAGAAA 1200 23 ω

AAF57997 standard; DNA; 24 쁑

AAF57997

0

26-APR-2001 (first entry)

Nucleic acid triplex DNA sequence

Hoogsteen-paired duplex; Watson-Crick pairing; triplex; antisense therapy; gene expression control; transcription;

Synthetic

WO200105937-A2

25-JAN-2001

20-JUL-2000; 2000WO-US019783.

20-JUL-1999; 99US-00357424. 19-JAN-2000; 2000US-00487130.

(TEXA) UNIV TEXAS.

DM, Hashem GM

WPI; 2001-159523/16.

Generating nucleic acid molecule comprising Hoogsteen-paired RNAasteriskDNA pyrimidinessteriskpurine duplex for use as an molecule, by heating a triplex to dissociate a Watson-Crick pyrimidine strand. paired

Example 1; Page 9; 23pp; English.

RESULT 270
AAP57997
ID AAP579
XX AAP579
XX AAP579
XX HOOGS
XX HOOGS
XX HOOGS
XX HOOGS
XX WO200
XX AAP51
XX WO200
XX AAP57
XX AAP5 The present invention describes a method for producing a nucleic molecule comprising a Hoogsteen-paired RNA:NNA duplex capable of medeaule comprising an RNA stoform a triplex. This involves Watson-Crick pairing. This is use antisense therapy, as it controls gene expression by causing transcription to be prevented. The present sequence is an example sequence used ń the methods example RNA sequence acid being о f

Sequence 24 BP; 12 A; 0 C; 12 G; 0 T; 0 U; 0 Other;

Query Match Best Local Similarity 0.3%; Score 17.8; DB 1; Pred. No. 4.1e+02; DB 1; Length

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RESULT 271
AAP57998/c
ID AAP57998/c
ID CAP57998/c
ID CAP57
RESULT 272
AAF57999/c
ID AAF579
XX
AC Nuclei
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Best Local S
Matches 19
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                            Nucleic acid triplex RNA sequence #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Generating nucleic acid molecule comprising Hoogsteen-paired RNAasteriskDNA pyrimidineasteriskpurine duplex for use as an antisense molecule, by heating a triplex to dissociate a Watson-Crick paired
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19-JAN-2000; 2000US-00487130.
                                                                                  26-APR-2001
                                                                                                                                                                                              AAF57999 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 24 BP; 0 A; 12 C; 0 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 9; 23pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleic acid triplex DNA sequence #3.
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Pred. No. 4
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Matches 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Generating nucleic acid molecule comprising Hoogsteen-paired RNAasteriskDNA pyrimidineasteriskpurine duplex for use as an antisense molecule, by heating a triplex to dissociate a Watson-Crick paired pyrimidine strand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hoogsteen-paired duplex; Watson-Crick pairing; triplex; antisense therapy; gene expression control; transcription; ss.
Nucleic acid triplex DNA sequence
                                                       AAF58000;
                                                                                                                                                                                                                                                                          Sequence 24 BP; 0 A; 12 C; 0 G; 0 T; 12 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 14; Page 17; 23pp; English
                            26-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-159523/16.
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                                                                                     AAF58000 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (TEXA ) UNIV TEXAS.
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19-JAN-2000;
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                                                                                                                                                          24
                                                                                                                                                                                                                   19;
                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-00357424.
2000US-00487130.
                            (first entry)
                                                                                   DNA;
                                                                                                                                                                                                                                90.5%;
                                                                                     24
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Pred. No. 4.1e+(
                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                4.1e+02
                                                                                                                                                                                                                                             Length 24;
                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                               an RNA sequence
                                                                                                                                                                                                                   0;
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AAF58000
ID AAF58000 standard; DNA; 24 BP.

XX
AC AAF58000;
XX
AC AAF58000;
XX
DT 26-APR-2001 (first entry)
XX
Nucleic acid triplex DNA sequence #4.

XX
Hoogsteen-paired duplex; Watson-Crick pairing; triplex;
XX
antisense therapy; gene expression control; transcription;
XX
OS Synthetic.
XX
PN W0200105937-A2.
XX
PD 25-JAN-2001.
XX
25-JAN-2000; 2000WO-US019783.
XX
20-JUL-1999; 99US-00357424.
PR 19-JAN-2000; 2000WS-00487130.
XX
TEXA ) UNIV TEXAS.
XX
Gray DM, Hashem GM;
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RESULT 274
AAF57996/c
ID AAF57996/c
ID AAF579
XX AAF579
XX AAF579
XX Nuclei
XX Hoogst
KW Hoogst
KW ANTise
XX Synthe
XX Synthe
XX 20-JUI
PD 25-JAN
XX 20-JUI
PR 19-JAN
XX 20-JUI
PR 20-JUI
XX Examp
XX Genera
PT RNAass
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Matches 19
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                                                                          The present invention describes a method for producing a nucleic acid molecule comprising a Hoogsteen-paired RNA: DNA duplex capable of being used as an antisense molecule and capable of recognising an RNA sequence to form a triplex. This involves Watson-Crick pairing. This is useful in
                                                                                                                                                                                                                                                                                       Generating nucleic acid molecule comprising Hoogsteen-paired RNAasteriskDNA pyrimidineasteriskpurine duplex for use as an molecule, by heating a triplex to dissociate a Watson-Crick projectie, by heating a triplex to dissociate a Watson-Crick projectie.
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  antisense therapy, as it controls gene expression by transcription to be prevented. The present sequence i triplex sequence used in the methods of the invention
                                                                                                                                                                                                                  Example
                                                                                                                                                                                                                                                                    pyrimidine
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19-JAN-2000; 2000US-00487130.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           therapy; gene
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                                                                                                                                                                                                                  Page
                                                                                                                                                                                                                                                                       strand.
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                                                                                                                                                                                                               9; 23pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 expression control;
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Pred. No. 4
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                                 sequence is
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RESULT 276
AAF87781
ID AAF877
XX
AC AAF877

standard;

mRNA;

24

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ARESULT 275
AAF5800/C
ID AAF580
XX AAF580
XX AAF580
XX AAF580
XX CAAFR
XX Nuclei
XX Hoogst
KW Hoogst
KW Antise
XX Synthe
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Best Local S
Matches 19
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                 The present invention describes a method for producing a nucleic acid molecule comprising a Hoogsteen-paired RNA.DNA duplex capable of being used as an antisense molecule and capable. The recognising an RNA sequence to form a triplex. This involves Watson-Crick pairing. This is useful in antisense therapy, as it controls gene expression by causing transcription to be prevented. The present sequence is an example of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Generating nucleic acid molecule comprising Hoogsteen-paired RNAasteriskDNA pyrimidineasteriskpurine duplex for use as an molecule, by heating a triplex to dissociate a Watson-Crick [
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hoogsteen-paired duplex; Watson-Crick pairing; triplex; antisense therapy; gene expression control; transcription;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleic acid triplex RNA sequence
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                                                                                                                                                                               Sequence
                                                                                                                                                                                                                             triplex sequence used in the methods
                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                pyrimidine strand.
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19-JAN-2000; 2000US-00487130.
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                                                                                                              Similarity
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                            AGAGAAAGAGAGAGAGAAA 1200
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  AGAGAGAGAGAGAGAGA
                                                                                                                                                                                                                                                                                                                                                                                                                  Page 17; 23pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hashem GM;
                                                                                                                                                                                  BP; 0
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                                                                                         Conservative
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Pred.
                                                                                                            Score 17.8;
Pred. No. 4.
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No. 4
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RESULT 277
AAF87784
ID AAF877
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AC AAF877
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AC AAF877
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DT 11-JUL
XX
DE Hybrid
XX
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                                                                                                                                                                                                                                                                                                                                                         The present invention describes a method for the identification of a CC nuclaic acid having a sequence capable of targeting a gene of interest CC comprises: (a) a first database having a list of stability values for CC independent combinations of N(x); (b) a computing unit having a means for CC inputting data comprising N(x), data list, defining a nucleic acid GC sequence of interest to be targeted to provide a second database; and (c) CC aprogram capable of processing the first and second database to N(x) (c) a program capable of processing the stability value of a nucleic acid sequence capable of comparison, and a stability value of a nucleic acid sequence capable of comparison, and a stability value of a nucleic acid sequence capable of comparison, and a stability value of a nucleic acid sequence of interest. The method having a sequence capable of targeting a gene of interest. The CC method may be used to obtain thermodynamic parameters for 20 combinations of nearest-neighbour base pairs of DNA;RNA hybrid sequences. The Nearest-CC Neighbour Thermal Stability Program can process data for use in CC calculating thermal melting temperatures for phosphorothicate DNA;RNA CC hybrids. The program can be readily extended to predict the most stable triplex-forming sequences, or antigene oligomers. The present sequence of the present sequence of the present sequence of the present sequence.
                                                                                                                                                                                                                                                                                    Best Loc
Matches
                                                                                                                                                                                                                                                                                                                     Query Match
                  Hybrid DNA
                                                                                                                                                                                                                                                                                                                                                      Sequence 24 BP; 12 A; 0 C; 12 G; 0 T; 0 U; 0 Other;
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03-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense DNA oligomer; ASO; identification; Nearest-Neighbour Thermal Stability Program; phosphorothioate; disease treatment; DNA:RNA
                                                      11-JUL-2001
                                                                                       AAF87784;
                                                                                                                          AAF87784 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Col 14; 43pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying a nucleic acid having a sequence capable of targeting a gene of interest, for identifying nucleic acids for gene therapy, comprises using the Nearest-Neighbor Thermal Stability Program.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (TEXA ) UNIV TEXAS SYSTEM.
                                                                                                                                                                                                                                                                                                      Local
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                                                                                                                                                                                                                                                                                                    Similarity
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                  sequence
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                                                     (first entry)
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97US-00808474.
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                                                                                                                          DNA;
                    SEQ ID NO:11
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thermal melting temperature;
hybrid; ss.
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Antisense DNA oligomer; ASO; identification; gene therapy; target; Nearest-Neighbour Thermal Stability Program; thermal melting temperature; phosphorothioate; disease treatment; DNA:RNA hybrid; ss.
                                                07-OCT-1994;
03-MAR-1997;
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                                                                                                                                                                                                                                                          Synthetic
(TEXA )
                                                                                                                     22-JAN-1999;
  UNIV TEXAS
                                                                     94US-00320507.
                                                  97US-00808474.
                                                                                                                     99US-00235614.
       SYSTEM.
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Identifying a nucleic acid having a sequence capable of targeting a ges of interest, for identifying nucleic acids for gene therapy, comprises using the Nearest-Neighbor Thermal Stability Program.

Gray DM,

Clark CL;

2001-280429/29.

Disclosure; Col 15; 43pp; English.

cc independent combinations of N(x); (b) a computing unit having a means for computing data comprising N(x), data list, defining a nucleic acid comparison of interest to be targeted to provide a second database; and (c) a program capable of processing the first and second database; and (c) comparison, and a stability value of a nucleic acid sequence capable of cargeting the gene of interest. The method is useful for identifying a nucleic acid having a sequence capable of targeting a gene of interest. The method may be useful to obtain thermodynamic parameters for 20 combinations of nearest neighbour base pairs of DN:RNA hybrid sequences. The Nearest of nearest neighbour base pairs of DN:RNA hybrid sequences. The Nearest calculating thermal Stability Program can process data for use in chybrids. The program can be readily extended to predict the most stable typics. Forming sequences, or antigene oligomers. The present sequence crepresents a hybrid DNA sequence which is used in the exemplification of The present invention describes a method for the identification of a nucleic acid having a sequence capable of targeting a gene of interest comprises: (a) a first database having a list of stability values for invention

Sequence 24 BP; 12 A; 0 C; 12 ဂ္ 0 7 0 U; 0 Other;

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                  Query Match
Best Local S
Matches 19
    1180
                   19;
                          Similarity
AGAGAAAGAGAGAGAGAAA 1200
                   Conservative
                         90.5%;
                  0;
                          Score 17.8;
Pred. No. 4.
                    Mismatches
                 4.1e+02;
2;
                                  DB
                                 1; Length 24;
                    Indels
                   0,
                  Gaps
                    0
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RESULT 278
AAF87783/c
ID AAF877
XX
AC AAF877
XX
DT 11-JUI
DX Hybric
XX
W Antisc
XW Neare
XW Neare
XW phospl
XX
OS Synth
                                                                                              AAF87783;
                                                                                                                  AAF87783 standard; mRNA; 24
                                                                           11-JUL-2001
                                                                            (first entry)
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Hybrid mRNA

sequence SEQ ID NO:10.

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AGAGAGAGAGAGAGAGA

22

Antisense DNA oligomer; ASO; identification; Nearest-Neighbour Thermal Stability Program; phosphorothioate; disease treatment; DNA:RNA gene therapy; target; thermal melting temperature; hybrid; ss.

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RESULT 279
AAF87782/c
ID AAF87782 standard; D
XX AAF87782;
AC AAF87782;
XX
DT 11-JUL-2001 (first
XX
DE Hybrid DNA sequence
XX
Antisense DNA oligom
XW Nearest-Neighbour Th
XW phosphorothicate; di
XX
OS Synthetic.
XX
PD US6183966-B1.
XX
PD 06-FEB-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC nucleic acid having a sequence capable of targeting a gene of interest comparises: (a) a first database having a list of stability values for comprises: (a) a first database having a list of stability values for comparises: (a) a first database having a list of stability values for computing data comprising N(x), data list, defining a nucleic acid comparison, and a stability value of a provide a second database; and (c) comparison, and a stability value of a nucleic acid sequence capable of comparison, and a stability value of a nucleic acid sequence capable of cargeting the gene of interest. The method is useful for identifying a concleic acid having a sequence capable of targeting a gene of interest. The method database treatment. The method may be used to obtain thermodynamic parameters for 2 combinations of nearest-neighbour base pairs of DNA:RNA hybrid sequences. The Nearest-CC Neighbour Thermal Stability Program can process data for use in calculating thermal stability Program can process data for use in calculating thermal melting temperatures for phosphorothicate DNA:RNA (C) hybrids. The program can be readily extended to predict the most stable triplex-forming sequences, or antigene oligomers. The present sequence content of the present sequence contents a hybrid many sequence which is used in the exemplification of the present sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 19
                                                                                                                                           Antisense DNA oligomer; ASO; identification; Nearest-Neighbour Thermal Stability Program; phosphorothicate; disease treatment; DNA:RNA
                                                                                                                                                                                                                          Hybrid DNA sequence SEQ ID NO:9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 24 BP; 0 A; 12 C; 0 G; 0 T; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes a method for the identification of a nucleic acid having a sequence capable of targeting a gene of interest
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Col 15; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gray DM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      07-OCT-1994;
03-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying a nucleic acid having a sequence capable of targeting a gene of interest, for identifying nucleic acids for gene therapy, comprises using the Nearest-Neighbor Thermal Stability Program.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US6183966-B1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1180 AGAGAAAGAGAGAGAGAAA 1200
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                                                                                                                                                                                                                                                                                                                                                                                                                                             23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Clark CL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94US-00320507.
97US-00808474.
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                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
                                                                                                                                                                                                                                                                                                                                               24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17.8;
Pred. No. 4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            No. 4.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 24;
                                                                                                                                         thermal melting hybrid; ss.
                                                                                                                                                                gene therapy; target; thermal melting tempe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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AAQ33955
AAQ33955
ID AAQ2
XX
AC AAQ2
AC AAQ2
AC Sequ
DT 02-F
DT 02-F
DX Sequ
XX
PCR,
KW PCR,
KW PCR,
KW Gen
XX
W092
XX
W092
XX
VX
PD 06-P
XX
PF 15-J
XX
PR 15-J
XX
PA (GEN

15-JAN-1991; 15-JAN-1992; 06-AUG-1992 WO9213102-A1 Bos taurus. genetic mapping;

9108-00642342 92WO-US000340

(GENM-) GENMARK

25-MAR-2003 02-FEB-1993

(revised) (first entry)

AAQ33955;

AAQ33955 standard;

DNA;

25 ВÞ

0

Sequence upstream from a microsatellite from clone TGLA351.

selection; primers; OPTIPRIM; breeding; cattle; parentage; tic mapping; traits; amplification; ss.

RESULT 280

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                                                                                                                                                                  CC nucleic acid having a sequence capable of targeting a gene of interest comprises: (a) a first database having a list of stability values for cindependent combinations of N(x); (b) a computing unit having a means for cindependent combinations of N(x); (b) a computing unit having a means for cindependent combinations of N(x); (b) a computing unit having a means for cindependent comprising N(x); (b) a computing unit having a means for computing data comprising N(x); (b) a computing unit having a means for computing the interest into the second database; and (c) comparison, and a stability value of a nucleic acid sequence capable of comparison, and a stability value of a nucleic acid sequence capable of targeting the gene of interest. The method is useful for identifying a concleic acid having a sequence capable of targeting a gene of interest. The method day are useful in gene therapy and disease treatment. The method may be used to obtain thermodynamic parameters for 20 combinations confidence thermal stability Program can process data for use in calculating thermal melting temperatures for phosphorothicate DNA:RNA hybrids. The program can be readily extended to predict the most stable triplex-forming sequences, or antigene oligomers. The present sequence crepresents a hybrid DNA sequence which is used in the exemplification of the present invention
                                                                                                   Query Match
Best Local (
                                                                                  Matches
                                                                                                                                                              Sequence 24 BP; 0 A; 12 C; 0 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identifying a nucleic acid having a sequence capable of targeting a gene of interest, for identifying nucleic acids for gene therapy, comprises using the Nearest-Neighbor Thermal Stability Program.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Col 14; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-280429/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-OCT-1994;
03-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TEXA ) UNIV TEXAS SYSTEM.
                          1180 AGAGAAAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ŋ.
24
                                                                                19;
                                                                                                   Similarity
  AGAGAGAGAGAGAGAGA 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Clark CL;
                                                                                                                                                                                                             invention
                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94US-00320507.
97US-00808474.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99US-00235614.
                                                                                                 90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     describes a method for
                                                                           0;
                                                                                                     Score 17.8; DB 1;
Pred. No. 4.1e+02;
                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          the
                                                                                                                       Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       identification
                                                                                  Indels
                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of a
                                                                                  Gaps
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RESULT 281
AAQ33606
ID AAQ336
XX AAQ336
XX AAQ336
XX AAQ336
XX AAQ336
DT 25-MAJ
DT 02-FE
XX MICTO
XX MCC
XX PCR;
KW Genet
XX Genet
XX W92
XX W992
XX W992
XX H5-;
XX 15-;
XX 15-;
XX H5-;
XX Gec
XX Ge
PA (GE
XX GE
XX PP
PA (GE
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequence is upstream of a bovine microsatellite sequence obtd. by CC screening a library of bovine MboI DNA fragments of between 250 and 500 bp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 CC clones cross-hybridised. Assuming independent distribution of microsatellites and MboI sites, the frequency of (T6)n >9 microsatellites in the bovine genome is estimated at >100, 000. The sequence information CC in the bovine microsatellites is summarised in the specification and indexed herein (see below). The sequences upstream and CC downstream of the microsatellite sequence were used to generate the crequired PCR primers for in vitro amplification of the corresp.

CC microsatellite (using the program OPTIFRIM). The microsatellites may be used to identify individuals, for parentage testing, and in the genetic mapping of economic trait loci, or genes involved the determinism of economically important traits esp. in cattle, to allow selective for in cattle, to allow selective considered PCR seeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         genetic mapping;
                                                                                                              Polymorphic bovine DNA
                                                                                                                                                                      WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                       15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                             06-AUG-1992.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9213102-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Microsatellite sequence from clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAQ33606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAQ33606 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Table
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Georges M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic bovine DNA markers - used in genetic identification, gene
                                                                                                                                                                                                                                                                                                                                  15-JAN-1991;
                                                                                                                                                                                                                                                                               (GENM-) GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     selection; primers; OPTIPRIM; breeding; cattle; parentage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTCATCTGGTGGCCACAGCAG 5089
                                                                                       and
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                            Page 172; 517pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Page 312;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CACATCTGGTGGCCACATCAG 22
                                                                                                                                                                                                                      Massey JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Маввеу JM;
                                                                               bovine DNA markers - used in genetic identification, selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               selective breeding.
                                                                                                                                                                                                                                                                                                                                     91US-00642342
                                                                                                                                                                                                                                                                                                                                                                                       92WO-US000340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           traits;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 7 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₽₽.
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Pred.
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No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGLA298.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .2e+02
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                                                                                                              gene
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RESULT 282
ACH61516/c
ID ACH615
XX
ACH615
ACH615
XX
ACH615

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation; library screening; Southern hybridisation; northern hybridisation; dot-blot hybridisation; gene sequence; mutation detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA target sequence #10652 useful in array for genetic analyses.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ACH61516 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACH61516;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (/TTIM)
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Pred. No. 4.
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The present invention relates to nucleic acid sequences that are complementary to particular genes, and can be used as probes for a variety of analyses such as gene expression analysis. Each probe comprises 9 or more consecutive nucleotides from at least one of 14936 nucleotide sequences defined in the patent, or their perfect sense match, sense mismatch, antisense match or antisense mismatch oligonucleotides. The probes may be used in an array comprising at least 10 distinct

New probe array useful e.g. for monitoring gene expression levels, analyzing genetic variations, or for hybridizing tag-labeled compoundables multiple nucleic acid probes.

compounds,

2003-576608/54

Claim 1;

SEQ ID

NO 10652; 9pp; English.

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RESULT 283
ACF79235/c
ID ACF792
XX ACF792
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Best Local S
Matches 19
                                                                                                            New calix(4)arene-nucleoside hybrid useful in gene therapy has one nucleoside attached to a calix(4)arene group through amide and is derived from a calix(4)arene having amino groups.
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/mod_base= OTHER
/note= "OTHER= ca
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                                                                           16pp; English.
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bonding,
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The present sequence is that of a calix(4)arene-oligonucleotide hybrid the invention, which includes a calix(4)arene-nucleoside (preferably thymidine) derivative. The calix(4)arene-oligonucleotide hybrid function

of.

Query Match

Sequence

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BP; 10 A; 13 C;

14 G;

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Other;

Score 17.8;

B

Length 50;

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                             The invention relates to a system for detecting gene expression, which comprises one or two isolated DNA molecules that detect expression of a gene, where the gene corresponds to any of 8143 oligonucleotides (ABZ00010-ABZ08152) each having 50 base pairs (bp). The system is useful for leukocyte expression profiling. It is particularly useful for disgnosing a disease, monitoring (rate of) progression of a disease, predicting therapautic outcome, determining prognosis for a patient, predicting disease complications in an individual or monitoring response to treatment in an individual. The diseases include cardiac allograft rejection, kidney allograft rejection, liver allograft rejection, atherosclerosis, congestive heart failure, systemic lupus erythematosus, rheumatoid arthritis, osteoarthritis or cytomegalovirus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             as a DNA hairpin structure mimic. It effectively recognises DNA or RNA through triplex formation by bonding between the calix(4) arene-containing cavity and a biologically active substance. The hybrid has a certain level of both rigidity and flexibility, is stable in vivo, has high cell permeability and can be mass-produced. It can be used as a DNA sensor or for gene therapy
                                                                                                                                                                                                                                                                               New system for leukocyte expression profiling, diagnosing a disease, or monitoring (the rate of) progression of a disease, e.g. atherosclerosis or congestive heart failure, comprises diagnostic oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 T7; leukocyte; gene expression profiling; allograft rejection;
atherosclerosis; congestive heart failure; systemic lupus erythematosus;
rheumatoid arthritis; osteoarthritis; cytomegalovirus; infection; probe;
                                                                                                                                                                                                                                               Claim 1; Page 341; Opp; English.
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Ly N, Woodward
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08-JUN-2001; 2001US-0296764P.
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d R, Quertermous T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA from an herbicide succeptible plant and an herbicide resistant plant of the same species, performing genetic analysis and identifying differences between their genomic DNA, identifying the difference that correlate with herbicide resistance or susceptibility, thus identifying polymorphic markers. The method is useful for identifying polymorphic markers of herbicide resistance in a plant e.g. Conyza canadensis, Lolium rigidum and goosegrass species, where the herbicides include glyphosate, paraquat and sulfonyl urea moieties. This sequence represents a primer associated with the identification of polymorphic markers of herbicide resistance
                                                                               AAV04338 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention describes a method of identifying polymorphic markers of herbicide resistance in a plant. The method involves: isolating genominations are considered to the control of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 6; Page 38; 168pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying polymorphic markers of herbicide resistance in a plant, by analyzing genomic DNA of herbicide resistant and susceptible plants, and identifying difference that correlate with resistance or susceptibility.
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Pred. No. 4
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence is a primer, which was used in the preparation of osteopyrotegerin (OPG) products. Anti-OPG antibodies can be used in OPG diagnostic assays, and as affinity purification materials. The OPG cDNA can be used to express recombinant OPG and to generate transgenic animals. It can also be used to regulate the level of OPG in mammals, specifically to increase OPG levels, however the use of antisense sequences is also contemplated. Fragments of the cDNA can be used as probes to detect OPG expressing cells and tissue, and to screen cDNA libraries for related sequences. OPG can be used to treat or prevent bor diseases, hypercalcaemia, hyperparathyroidism, rheumatoid arthritis, osteolytic metastases, periodontal bone loss, bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        osteoporosis; Paget's disease; hypercalcaemia; hyperparathyroidism; rheumatoid arthritis; osteomyelitis; osteolytic metastasis; periodontal bone loss; bone necrosis; osteopaenia; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Osteoprotegerin; antibody; diagnosis; affinity purification; recombinant production; transgenic animal; treatment; prevention; antiense oligonucleotide; probe; detection; screening; bone disease;
            Human; corticoliberin 26; cytostatic; virucide; immunodulatory; antiinflammatory; haemostatic; gene therapy; malignant tumour; haemopathy; HIV infection; immunological disease; inflammation;
                                                                                                                                                                                                                                                                                                                                                                      Sequence
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03-SEP-1996;
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                                                                       Human corticoliberin 26 coding sequence
                                                                                                                                      AAH73990;
                                                                                                                                                                    AAH73990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 5; Page 18; 182pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid encoding osteoprotegerin - useful for treatment of diseases involving excessive bone loss, e.g. osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-334271/31
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                                                                                                                                                                                                                                                                                                                                                                                                   necrosis and osteopaenia
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                                                                                                       (first
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Pred. No. 4.4e+02;
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                                                                         PCR primer
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Best Local S
Matches 20
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                                                                                                                                                                                                                                     Human; FD 17; cytostatic; virucidal; immunomodulatory; haemostatic; antiinflammatory; gene therapy; malignant tumour; haemopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human corticoliberin 26 and encoded polynucleotide, used in diagnosis treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
                                                                                              02-MAR-2000; 2000CN-00111868
                                                                                                                    26-FEB-2001; 2001WO-CN000221
                                                                                                                                                                  WO200164729-A1
                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                             immunological
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       example from the present invention
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 New human polypeptide FD
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                         WPI; 2001-550164/61.
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Pred. No. 4
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  and treating malignant tumor,
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Best Local S
Matches 20
 The present invention relates to a method for treating conditions leading to bone loss. The method comprises administering a purified and isolated osteoprotegerin (OPG) protein (ARE57836-ARE57838 and ARB66974-ARB66976) in conjunction with other substances such as tumour necrosis factor-alpha (TNF-alpha) inhibitors, interleukin (IL)-6, -8 and -18 inhibitors, ICE modulators, fibroblast growth factor (FGF)1-10 modulators and/or platelet
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bone loss; osteoprotegerin; OPG; rheumatoid arthritis; hyperalgesia; multiple sclerosis; osteoproteger osteoprotegerin; osteoproteitis; asthma; inflammation; studies of the sclerosis; graft-versus-host disease; septic shock; acute pancreatitis; Alzheimer's disease; anorexia; atherosclerosis; pacoronary condition; myocardial infarction; cancer; diabetes; psoriasis endometriosis; fever; glomerulonephritis; inflammatory bowel disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes the human FD 17 protein (I). (I) has cytostatic, virucidal, immunomodulatory, antiinflammatory and haemostatic cytostatic. The polymucleotide encoding (I) can be used in gene therapy. (I) and the polymucleotide encoding it are applicable in the diagnosis and treatment of malignant tumour, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. The present sequence represents a PCR primer for human FD 17, which is
                                                                                                                                                Treating conditions leading to bone loss such as rheumatoid arthritis, multiple sclerosis and asthma, comprises administering an osteoprotegerin protein in conjunction with e.g. inhibitors of interleukin and tumor
                                                                                                                                                                                                                                                                                                           09-JUL-1999;
09-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     hemopathy, human immunodeficiency virus (HIV) infection, diseases and inflammations.
                                                                                                         Example
                                                                                                                                                                                                                                                Boyle WJ,
                                                                                                                                                                                                                                                                                                                                                       07-JUL-2000; 2000WO-US018667.
                                                                                                                                                                                                                                                                                                                                                                                       18-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                     WO200103719-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human OPG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF57740;
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                                                                                                                                      necrosis factor alpha.
                                                                                                                                                                                                                                                                               (AMGE-) AMGEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ischaemia; Parkinson's disease; PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   an example from the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR
                                                                                                                                                                                                                                                Lacey DL,
                                                                                                        Page 88; 316pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 11; 36pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 0 A; 2 C; 1 G; 21 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                               INC.
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                                                                                                                                                                                                                                                                                                           99US-00350670.
99US-00457647.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
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Pred. No. 4.4e
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disease;
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fibroblast growth

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RESULT 290
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Matches 20
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                                                                                                                                  The invention relates to an isolated polypeptide of human tre carcinogenic gene protein 10.56 comprising a 96 residue amino acid sequence, fully defined in the specification, or its fragment, analog or derivative. The polypeptide is useful in the diagnosis and treatment of malignant tumours, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammatory disorders present sequence is a primer used to amplify a polynucleotide encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human tre carcinogenic gene protein 10.56 and encoding polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; Tre carcinogenic gene protein 10.56; cytostatic; haemostatic; virucide; immunomodulatory; antiinflammatory; gene therapy; cancer; haemopathy; human immunodeficiency virus; HIV; infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Page 17; 36pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-083078/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-MAY-2000; 2000CN-00115824.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SHAN-) SHANGHAI BIOWINDOW GENE
                                                                                                polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               carcinogenic gene protein 10.56 PCR primer 2.
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                                                                                                     invention
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immunological diseases
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                                                                                                                                  s inflammatory disorders. The a polynucleotide encoding
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                                        The invention relates to the use of phytanic acid or its derivatives for the treatment or prevention of diabetes mellitus. The invention also relates to a method for treating or preventing non-insulin dependent diabetes mellitus (NIDM) or other conditions associated with impaired glucose tolerance such as obesity using phytanic acid or its derivatives. The phytanic acid, their derivatives or their precursors are useful as pharmaceutical compounds or supplements to foods or feeds for the treatment or prevention of type II or NIDDM, hyperlipidaemia, hyperrocolesterolaemia, hyperinsulinaemia, syndrome X, hypertension, hypertriglyceridaemia, impaired glucose tolerance and related obesity. They are also useful in insulin therapy in combination with known active compounds. The present sequence is glucokinase cDNA specific reverse transcription PCR (RT-PCR) probe used in the exemplification of the
                                                                                                                                                                                                                                                                                                   New composition comprising phytanic acid or its derivatives, useful for treating or preventing non-insulin dependent diabetes mellitus, impaired glucose tolerance and related obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-AUG-2000; 2000EP-00116848
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RT-PCR; ss.
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ABA99264/c
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                recombinant production and antagonist, encoding polynucleotide and application. The polypeptide is useful for treating primary T lymphocyte immune deficiency disease, human immunodeficiency virus infection, other immune diseases, various tumours, embryonic development disturbance diseases, growth and development obstruction disease, inflammation and haemopathy. The present sequence is that of a PCR primer, useful to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; immune response factor 64; infection; HIV; inflammation; primary T lymphocyte immune deficiency disease; haemopathy; tumour; human immunodeficiency virus; immune diseases; growth; PCR primer; embryonic development disturbance; development obstruction disease.
                                ABA99264;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP, 11 A; 2 C; 5 G; 6 T; 0 U; 0 Other;
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                                                                                        ABA99264 standard;
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RESULT 294
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Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel human tre oncogene 10.56 which has cytostatic, haemostatic, virucide, immunomodulatory and antiinflammatory activity and can be used for gene therapy. The polypeptide of the invention and its encoding polynucleotide are used in diagnosis and treatment of malignant tumours, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammations. This sequence represents an RT-PCR primer used in the amplification of the human tre oncogene 10.56 gene which is described in the disclosure of the
                                                                                                                                             compound activity characterisation; cellular activity; phenotypic attribute; candidate medicine; candidate tr multiple biological descriptor; cell marker; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oncogene; tra oncogene 10.56; human; treatment; gene therapy; cytostatic; haemostatic; virucide; immunomodulatrory; antiinflammatory; diagnosis; malignant tumour; haemopathy; human immunodeficiency virus; HIV infection; immunological disease; inflammation; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 human tre oncogéne 10.56 and encoding polynucleotide, used in diagno and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and inflammation.
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       24-APR-2001; 2001WO-US013248.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 compound activity characterisation; cellular activity; phenotypic attribute; candidate medicine; candidate tr multiple biological descriptor; cell marker; ss.
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                                                                                          Oestreicher DR,
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20-FEB-2001; 2001US-00790214.
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20-FEB-2001;
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                                                                                          Sabry JH,
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Pred. No. 4
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                                                                                          Vaisberg EA,
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RESULT 296
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Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Characterizing cellular activity of compound, by receiving images cells with known activity and images of cells treated with compound characterizing phenotypic attributes of images and comparing the
The invention relates to a novel zinc finger protein designated 11.55. Also disclosed are the polymucleotide encoding it, and a process for preparing the polymeptide using but the encodination techniques. The application of the polymeptide is in treating diseases such as cancer and human immunodeficiency virus (HIV) infection, The current sequence represents a zinc finger protein 11.55 related PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Fig 18; 139pp; English.
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                                                                                                                                                                                                                                                The present invention describes a method for identifying a pancreatic ductal carcinoma-specific gene. The method comprises preparing pancreatic ductal cells from a pancreatic ductal carcinoma patient and a normal individual. Also described: (1) a method of testing for pancreatic ductal carcinoma; (2) a method for identifying a drug candidate compound for treating or preventing pancreatic ductal carcinoma; and (3) a drug for testing for pancreatic ductal carcinoma comprising, as an active ingredient, a molecule selected from an antibody binding to a protein encoded by a pancreatic ductal carcinoma-specific gene and an oligonucleotide specifically hybridising to a transcription product of a pancreatic ductal carcinoma-specific gene or comprising a compound is useful in identifying a pancreatic ductal carcinoma-specific gene has cytostatic activity, and can be used in gene therapy. The method is useful preparing compounds for treating and/or preventing pancreatic ductal carcinoma. The present sequence represents a PCR primer which is
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nilarity 83.3%;
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                                                                                                                                   Matches
                                                                                                                                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                    chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal celicarcinoma. This oligonucleotide sequence is an immunostimulatory non-CpG variant DNA oligo, used in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New immunostimulatory oligonuclectide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
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                                                                                                                                                                                                                                                                          Sequence 24 BP; 1 A; 1 C; 1 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 14; SEQ ID NO 21; 139pp; English
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                                                                                                                                                                       Local
                                                                 5396 AAAATACAAAAAAGAAAAAATGAA 5419
                                                                                                                                   1 Similarity
20; Conserv
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83.3%;
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Pred. No. 4.
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RESULT 299 ADG75920/c ID ADG759

ADG75920 standard; DNA; 24 BP.

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RESULT 300
ADG75918/c
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AC ADG759
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Best Local S
Matches 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                        chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell
carcinoma. This oligonucleotide sequence is an immunostimulatory non-CpG
variant DNA oligo, used in an exemplification of the invention.
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plasmacytoid dendritic cell; immunomodulator; gene therapy;
chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
                    Immunostimulatory non-CpG oligonucleotide IMT 173 SeqID
                                                       11-MAR-2004
                                                                                          ADG75918
                                                                                                                           ADG75918 standard;
                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 1 A; 1 C; 1 G; 21 T; 0 U; 0 Other;
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ADM28709
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          contain a non-palindromic sequence motif. Specifically, it refers to DNA cligonucleotides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune response is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation and cytokine production in plasmacytoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an cantigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is an immunostimulatory non-CpG variant DNA oligo, used in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local
Human; OPG; ss; primer; bone resorption; excessive bone loss; osteoporosis; Paget's disease of bone; hypercalcaemia; hyperparathyroidism; steroid-induced osteopaenia; rheumatoid arthritis; hyperparathyroidism; steroid-induced osteopaenia; rheumatoid arthritis; osteomyelitis; osteolytic metastasis; periodontal bone loss; Cushing's syndrome; acromegaly; osteogenesis imporfecta; homocystinuria; Menke's syndrome; Riley-day syndrome; immobilisation of extremity; tumour; haematologic malignancy; multiple myeloma; lymphoma; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 24 BP; 1 A; 1 C; 1 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 14; SEQ ID NO 20; 139pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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                                                                                                                                                                                                                                                                              PCR primer #2.
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Pred. No. 4.
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Cystedine rich domains of tumour necrosis factor receptor extracellular color domains of tumour necrosis factor receptor extracellular color its fragment which specifically binds to OPG, a composition comprising CC OPG (in a carrier, adjuvant, solubiliser, stabiliser and/or anti-oxidant) and an osteoprotegerin multimer consisting of osteoprotegerin monomers. CC Ab is useful for detecting the presence of OPG in a biological sample complete incubating the sample with Ab under conditions that allow CC which involves incubating the sample with Ab under conditions that allow CC binding of ab to OPG and detecting the bound Ab. OPG is useful for regulating the levels of OPG in an animal (human). The nucleic cuseful for regulating the levels of OPG in an animal (human). The nucleic colored promotes an increasing in tissue level of OPG. OPG NA is cold promotes an increasing in tissue level of OPG. OPG is useful for colored promotes an increasing in tissue level of OPG. OPG is useful for colored promotes an increasing in tissue level of OPG. OPG NA is consequently a bone loss one loss due to osteopaenia, bone loss due to rheumatoid arthritis, bone loss due to osteopaenia, bone loss due to rheumatoid arthritis, bone loss due to osteopaenia protein BMP-1 through BMP-12, TGP-beta family members, IL-1 considered without and their analogues, and periodontal bone loss. The costeoporosis (hyperthyroidism, cushing's syndrome, and their analogues, and service of constaglandins, bisphosphonates, and bone-enhancing minerals. OPG is costeoporosis (hyperthyroidism, cushing's syndrome, and acromegaly).

CC hereditary and congenital forms of osteoporosis (osteogenesis imperfecta consequences) and their analogues, and acromegaly).

CC hereditary from solid tumours and haematologic malignancies (multiple costeopaenia associated with hyperthyroidism and renal function colorated with hyperthyroidism and renal function coll disorders, osteopaenia following surgery and osteopaerosis or bone cell death. The present sequences is an OPG cDNA cl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BOYL/) BOYLE W J.
(LACE/) LACEY D L.
(CALZ/) CALZONE F J.
(CHAN/) CHANG M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a purified and isolated polypeptide having osteoprotegerin (OPG), an OPG polypeptide from rat, human and mouse, or having amino terminus at residue 22, and 1-216 amino acids are deleted from carboxy terminus of human OPG polypeptide. Also included are an isolated nucleic acid encoding an OPG polypeptide (OPG NA), an expression vector comprising OPG NA, a host cell transformed or transfected with the vector, a transgenic mammal comprising the cell, producing OPG, a polypeptide comprising an amino acid sequence of at least about 164 amino acids comprising four cysteine-rich domains characteristic of the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Boyle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-NOV-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-DEC-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2003207827-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            osteoprotegerin; transgenic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 21; 141pp; English.
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96US-00706945.
96US-00771777.
98US-00132985.
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Best Local
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                                                                                                                                                                                                                                                                        The invention relates to a novel isolated nucleic acid encoding a Monarch -1, CATERPILIER (CARD [Caspase recruitment domain], transcription enhancer, R (purine)-binding, pyrin, lots of leucine repeats) 11.2, CATERPILIER 11.3, CATERPILIER 16.2 or CIAS1 (cold-induced autoinflammatory syndrome 1) polypeptide comprising the amino acids encoded by exon 6 and lacking the amino acids encoded by exon 6 and lacking the amino acids encoded by exon 4 or its fragment. The nucleic acid of the invention demonstrates antiinflammatory and cytostatic acityities and may be useful in preparing a composition for treating an inflammatory disease or cancer, possibly via gene therapy. The current sequence is that of a CATERPILER-related RT-PCR primer which was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acid encoding Monarch-1, CATERPILLER 11.2, CATERPILLER 11.3, CATERPILLER 16.1, CATERPILLER 16.2 or CIAS1 polypeptide, useful in preparing a composition for treating inflammatory disease or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CARD, transcription CATERPILLER 11.3; C
                                                                                                                                                                                                                      Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 2; SEQ ID NO 119; 205pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-348215/32
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cold-induced autoinflammatory syndrome 1; antiinflammatory; cytostatic;
inflammatory disease; cancer; gene therapy; human; purine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Monarch-1; CATERPILLER 11.2; caspase recruitment domain; CARD, transcription enhancer, R-binding, pyrin, lots of leucine repeat; CATERPILLER 11.3; CATERPILLER 16.1; CATERPILLER 16.2; CIAS1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CATERPILLER 19.3; chromosome 19q13; multiple sclerosis; PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RT-PCR primer 2 used to analyse human Monarch-1 expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYNC-) UNIV NORTH CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local
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                                                                                                                                          Similarity
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                                                    CTTCTACTTCAGCTGCAGAGAGCT 1637
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Davis B,
                                                                                                                                                                                                                         BP; 6 A;
                                                                                                            Conservative
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                                                                                                                                                                                                                         5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                    83.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    83.3%;
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                                                                                                                                 Score
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                                                                                                                                          No. 4
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No. 4.
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                                                                                                                                                                    DB 1;
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J, 2
                                                                                                                                                           Length
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ADP27725/c
ID ADP27725 standard;
                                                  RESULT 304
                                                                                                                                                                                                                                                                associated with EGFR (epidermal growth factor receptor) positive cancer. Specifically, it refers to a gene expression profiling method that can provide a prediction as to whether a patient is likely to respond well to treatment with an EGFR inhibitor. The present invention describes the quantitative analysis of the expression level of the RNA transcript of at least one gene selected from the group of CD44V3, CD44V6, DR5, GR01, KRT17, LAMC2 or their products thereof. It further provides a cDNA microarray containing named genes that represent prognostic transcripts which are useful for determining whether a patient diagnosed with an EGFR expressing head or neck cancer or colon cancer exhibits elevated or decreased expression levels of these genes compared to normal. As such, these methods are also useful for prognosing or predicting the likelihood of cancer-attributable death or progression, including recurrence and metastatic spread of a neoplastic disease, as well as drug resistance. This oligonucleotide sequence is a PCR primer used to amplify a human PCR amplicon DNA sequence used as a prognostic cancer marker, given in an exemplification of the invention.
                                                                                                                                                                   Query Match
Best Local S
Matches 20
                 AAX33267 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Prognosing a patient with EGFR-expressing colon cancer comprises subjecting a sample comprising EGFR-expressing cancer cells to quantitative analysis of the expression level of the RNA transcript
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                microarray; head and
neoplastic disease;
                                                                                                                                                                                                                                    Sequence 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-NOV-2003; 2003WO-US036777.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JUN-2004
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epidermal grow
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to a novel method concerning prognostic markers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-NOV-2002; 2002US-0427090P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer to amplify a human cancer prognostic marker DNA SeqID 162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-AUG-2004
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                                                                                                                                3257 AGGACCTGGCCTCTGTGCTTAGTG 3280
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               growth factor receptor; cancer; gene expression profiling;
y; head and neck cancer; colon cancer; metastatic spread;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 162; 113pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene e.g.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cronin MT,
                                                                                                                                                                                                                                  BP; 7
                                                                                                                                                                   0.3%;
llarity 83.3%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
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                                                                                                                                                                                                                                    A; 9 C; 5 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Shak S,
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                   ВP
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                                                                                                                                                                   Score 17.6; DB 1; Length 24; Pred. No. 4.4e+02; O; Mismatches 4; Indels
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                                                                                                                                                                  Gaps
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RESULT 305
AAC95663/C
ID AAC966
XX AAC966
XX AAC966
XX AAC966
XX DAAC966
XX DAAC966
XX USACAL HA HI
XX USACAL 
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Best Local Similarity
Matches 20; Conserv
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15-OCT-1997;
08-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes DNA which participates in the regulation of expression of PBBP2 alpha A gene. The DNA produces a regulator protein with the activity of promoting bone formation and can serve as a promote for prevention and treatment of bone diseases including osteoporosis. The present sequence represents a PCR primer used in an example from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA regulating expression of PEBP2 alphaA gene to protein, useful as promoter for prevention or/and diseases e.g. osteoporosis.
                                                02-NOV-2000
                                                                                                                                                                                                                      DNA sequence analysis; sequencing; protein sequence; protein structure; gene typing; organ donation; bacteria identification; 16s rRNA; HLA;
                                                                                                                                                                                                                                                                                                                                                                                                                                               AAC96663 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-JUN-1999
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                                                                                              WO200065088-A2
                                                                                                                                                                                             human leukocyte antigen; PCR primer; 88.
                                                                                                                                                                                                                                                                                                HLA HLA-A gene PCR primer #40.
                                                                                                                                                                                                                                                                                                                                                 26-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                 AAC96663;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 6; Page 39; 118pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nakateuka
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9911787-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (SUMU ) SUMITOMO PHARM CO LTD.
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                                                                                                                                                 sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                 (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97JP-00254250.
97JP-00299407.
98JP-00114135.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; 5 C; 4 G; 2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                                                                                 entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.6; DB 1;
Pred. No. 4.5e+02;
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treatment of bone
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20-APR-2000; 2000WO-EP003636.

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RESULT 306
ABN13298
ID ABN132
XX ABN132
XX ABN132
XX Human
XX Human
XX Human
XX Homo
XX WO200
XX WO200
XX WO200
XX For 25-MA
XX 26-MA
PR 21-SEE
PR 27-SEE
PR 27-SEE
PR 30-JJ
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Best Local
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                          26-MAY-2000;
21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                04-OCT-2000;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              muscle; myosin;
skeletal muscle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human GDMLP-1 25-mer scanning SEQ ID NO:5 sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention provides a method for identifying a set of extendible primers which can be used in the identification, typing and classification of genes. This can then be used to predict protein sequence and structure, in organ donation to match the organ with the receiver, and to identify bacteria in a sample. The method can be used type the human leukocyte antigen genes (HLA) and 16s rRNA genes in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Identifying extendible primers for use in identification, or classification of a nucleic acid of an organism, allele or gene such as class 1/2 HLA comprises identifying all possible nucleotide sequences of
                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200192524-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 25 BP; 2 A; 3 C; 2 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                               25-MAY-2001; 2001WO-US016981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; genome-derived myosin-like protein 1; GDMLP-1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABN13298 standard; DNA; 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5392 ТАЛАЛАЛТАСАЛАЛАДОЛАЛАЛ 5415
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                   ; 2000US-0207456P.
; 2000US-023458PP.
; 2000US-0236359P.
; 2001WO-US000661.
; 2001WO-US000662.
; 2001WO-US000664.
; 2001WO-US000665.
; 2001WO-US000665.
; 2001WO-US000665.
; 2001WO-US000667.
; 2001WO-US000667.
; 2001WO-US000668.
; 2001WO-US000668.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              chromosome 22; gene disorder; amplicon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17.6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              therapy; vaccine; heart disease; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .5e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO:13290.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25,
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CC protein 1 (hopMLP-1). The protein and polynucleotide sequences of hGDMLP-1 CC nucleic acids can be used as probes to detect, characterise and quantify CC hGDMLP-1 nucleic acids in samples, as amplification substrates, to CC protein variants shaving desired phenotypic improvements, and for CC expressing the proteins. The hGDMLP-1 proteins or polypeptides may be CC expressing the proteins. The hGDMLP-1 proteins or polypeptides may be CC used as immunogens to raise antibodies that specifically recognise hGDMLP-1 cc proteins, as standards in assays used to determine the concentration CC and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific biomolecule conduction, and in vaccines or for replacement therapy. The proteins and in vaccines or for replacement therapy. The conduction, and in vaccines or for replacement therapy. The conduction, and in vaccines or for replacement therapy. The conduction, and in vaccines or for replacement therapy. The conduction and in vaccines or for replacement therapy. The conduction and in vaccines or for replacement therapy. The conduction and in vaccines or for replacement therapy. The conduction muscle disorders. hGDMLP-1 may be used for diagnosing a conduction of the present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. CC hGDMLP-1 sequence are presents an oligomer used in the screening of the present data for this patent did not form part of the printed CC specification, but was obtained in electronic format directly from WIPO at the present sequence data for this patent did not form part of the printed CC specification.
  New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gu Y,
  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present protein 1 ()
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  BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     invention describes a human genome-derived myosin-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ID NO 13290;
A, 3 C; 11 G; 3 T; 0 U;
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     0 Other;
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Matches
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26-MAY-2000;
21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
30-JAN-2001;
                                                                                                                                     Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                                           25-MAY-2001, 2001WO-US016981.
                                                                              06-DEC-2001.
                                                                                                WO200192524-A2
                                                                                                                                                                             Human
                                                                                                                                                                                                29-MAY-2002
                                                                                                                                                                                                                                      ABN12957 standard;
                                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                                                                    3475 AGCAGACGGAAACCAAGTGTGATG 3498
                                                                                                                                                                            GDMLP-1 25-mer scanning
                                                                                                                                                                                                                                                                                      N
                                                                                                                                                                                                                                                                                                                            Similarity
20; Conserv
  ; 2000US-0207456P.
; 2000US-0234687P.
; 2000US-0236359P.
; 2000GB-00024263.
; 2001WO-US000661.
                                                                                                                                                                                                                                                                                                                            0.3%;
llarity 83.3%;
Conservative
                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                      25
                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                            Score 17.6; DB 1;
Pred. No. 4.5e+02;
0; Mismatches 4
                                                                                                                                                                             SEQ ID NO:5 sequence SEQ ID NO:12949.
                                                                                                                                                                                                                                                                                      25
                                                                                                                                                                                                                                                                                                                                              DB 1; Length 25;
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                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                hGDMLP-1; heart; heart disease;
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ARBSULT 308
ARBNO4840
ID ABNO4840
AC ABNO48
AC
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polymucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 2.
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30-JAN-2001;
30-JAN-2001;
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ည
                                                                                         Human; genome-derived myosin-like protein 1; GDMLP-1; muscle; myosin; chromosome 22; gene therapy; vaccine; skeletal muscle disorder; amplicon; screening; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1
                                                                                                                                                                                                               Human GDMLP-1
                                                                                                                                                                                                                                                                                                                                     ABN04840
                                                                                                                                                                                                                                                                                                                                                                                             ABN04840 standard; DNA; 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2112 GATGCAGCAGATGAAGCGGAAGGA 2135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 10 A; 4 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001WO-US000664.

2001WO-US000665.

2001WO-US000666.

2001WO-US000667.

2001WO-US000669.

2001WO-US000670.
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2001WO-US000663.
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                                                                                                                                                                                                            25-mer
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                                                                                                                                                                                                         scanning
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No.
                                                                                                                                                                                                            SEQ ID NO:4 sequence SEQ ID NO:4832.
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                                                                                                                        therapy; vaccine; heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                .5e+02
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                                                                                                                                                    hGDMLP-1; heart;
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RESULT 309
ABN04839
ID ABN048
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AC ABN048

ABN04839 standard; DNA; 25

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Best Local
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30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
30-JAN-2001;
                                                                                                                                                                                                                                                                                                                                 protein 1 (hGDMLP-1). The protein and polynuclectide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hGDMLP-1 protein variants having desired phenotypic improvements, and for expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP proteins, as standards in assays used to determine the concentration and/or amount specifically of hGDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser description ionisation, as
                                                                                                                                                           therapeutic supplement in patients having specific deficiency in hGDMLP-1 production, and in vaccines or for replacement therapy. The polynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart and skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22. The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
                                                                                                                                 Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gu Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200192524-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-SEP-2000;
27-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AEOM-) AEOMICA INC.
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                                                                                Local Similarity
                                 3257 AGGACCTGGCCTCTGTGCTTAGTG 3280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002-179446/23.
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2001WO-US000662.
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2001WO-US000664.
2001WO-US000664.
2001WO-US000666.
2001WO-US000666.
2001WO-US000666.
2001WO-US000666.
                                                                                                                               BP; 4 A; 8 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention
                                                                Conservative
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2001US-0266860P
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                                                                                0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4832; 214pp; English.
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                                                                0,
                                                                                Score 17.6;
Pred. No. 4.
                                                                  Mismatches
 24
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                                                                                  6; DB 1;
4.5e+02;
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                                                                                                Length 25;
                                                                  Indels
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                                                                  <u>,,</u>
                                                                Gaps
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29-MAY-2002

(first entry)

GDMLP-1 25-mer scanning

SEQ ID NO:4 sequence

SEQ

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NO:4831.

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CC protein 1 (hGDMLP-1). The protein and polymucleotide sequences of hGDMLP-1 CC nucleic acids can be used as probes to detect, characterise and quantify CC hGDMLP-1 nucleic acids in samples, as amplification substrates, to protein variants having desired phenotypic improvements, and for CC provide initial substrates for the recombinant engineering of hGDMLP-1 CC protein variants having desired phenotypic improvements, and for CC expressing the proteins. The hGDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hGDMLP-1 proteins, as specific blomolecule capture probes for surface-enhanced laser desorption ionisation, as capture probes for surface-enhanced laser desorption ionisation, as capture probes for surface-enhanced laser desorption ionisation, as continuous and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The copynucleotide sequences encoding hGDMLP-1 may be used for diagnosing a disorder associated with the expression of hGDMLP-1, in particular heart cand skeletal muscle disorders. hGDMLP-1 is localised to chromosome 22.

CC The present sequence represents an oligomer used in the screening of the hGDMLP-1 sequence in the exemplification of the present invention. N.B. CT has been accounted at a for this patent did not form part of the printed CC specification, but was obtained in electronic format directly from WIPO at figure of the present directly published pct_sequence
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21-SEP-2000;
27-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sent invention describes a human genome-derived myosin-like 1 (hGDMLP-1). The protein and polynucleotide sequences of i
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2000US-0236359P.
2000GB-00024263.
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G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       therapy;
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Query Match Best Local S Matches 20

20; Conservative

0.3%;

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Score 17.6; DB 1; Pred. No. 4.5e+02; 0; Mismatches 4

DB 1; Length 25;

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0 Other;

The present invention describes a human genome-derived myosin-like CC protein 1 (hgDMLP-1). The protein and polynucleotide sequences of hgDMLP-1 can be used in gene therapy and vaccine production. The hgDMLP-1 cC nucleic acids can be used as probes to detect, characterise and quantify concleic acids in samples, as amplification substrates, to provide initial substrates for the recombinant engineering of hgDMLP-1 cC protein variants having desired phenotypic improvements, and for expressing the proteins. The hgDMLP-1 proteins or polypeptides may be used as immunogens to raise antibodies that specifically recognise hgDMLP-1 proteins, as standards in assays used to determine the concentration and/or amount specifically of hgDMLP proteins, as specific biomolecule capture probes for surface-enhanced laser desorption ionisation, as complement in patients having specific deficiency in hgDMLP-1 production, and in vaccines or for replacement therapy. The production, and in vaccines or for replacement therapy. The constitute sequences according hgDMLP-1 may be used for diagnosing a constitute associated with the expression of hgDMLP-1 in particular heart cand skeletal muscle disorders. hgDMLP-1 is localised to chromosome 22.

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RESULT 310
ABN12958
ID ABN1295
XX ABN129
XX Human
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21-SEP-2000;
27-SEP-2000;
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RESULT 311
ABN13300
ID ABN133
XX ABN133
PR 25-MA1
XX 26-MA1
XX ABN133
PR 27-SEI
PR 27-SEI
PR 27-SEI
PR 30-JA1
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Best Local Similarity
Matches 20; Conserv
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21-SEP-2000;
27-SEP-2000;
04-OCT-2000;
                    The present invention describes a human genome-derived myosin-like protein 1 (hGDMLP-1). The protein and polynucleotide sequences of hGDMLP-1 can be used in gene therapy and vaccine production. The hGDMLP-1 nucleic acids can be used as probes to detect, characterise and quantify hGDMLP-1 nucleic acids in samples, as amplification substrates, to
                                                                                                                                                                                                                             New polypeptide, for raising antibodies that recognize hGDMLP-1 proteins, or as specific biomolecule capture probes for surface-enhanced laser desorption ionization, comprises human myosin-like protein hGDMLP-1.
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                                                                                                                                                                                 Disclosure;
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2000US-0234687P.
2000US-0236359P.
2000GB-00024263.
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Pred. No. 4.5e
0; Mismatches
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Best Local S
Matches 20
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30-JAN-2001; 2001WO-US000664.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000667.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000668.
30-JAN-2001; 2001WO-US000669.
23-MAY-2001; 2001WO-00064761.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; gene therapy; tumour suppressor; HTPL; chromosome 10p12.1; human testis expressed Patched like protein; testis; adrenal; liver; male germ cell development; bone marrow; brain; kidney; lung; placenta; prostate; skeletal muscle; colon; male infertility; cancer; ss.
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Pred. No. 4.
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Novel isolated human testis expressed Patched like protein (HTPL), for identifying agonist and antagonist and specific binding partner for treating subjects having defects in HTPL.

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Example

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Page 341; 718pp; invention

English

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testis expressed Patched like

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RESULT 313
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Matches 20
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This invention describes a novel isolated nucleic acid that encodes one of three new isoforms of human pregnancy associated plasma protein E, hPAPP-E. The products of the invention have abortive and contraceptive activity and can be used for gene therapy or in a vaccine. The nucleic acid, polypeptide encoded by it, or antibody to the polypeptide can be used in pharmaceutical compositions or vaccines for preventing or abouting pregnancy. PAPP-E is used in the antenatal diagnosis of

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New isolated nucleic acid

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Example associated

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                           This invention describes a novel isolated nucleic acid that encodes one of three new isoforms of human pregnancy associated plasma protein E, hPAPP-E. The products of the invention have abortive and contraceptive activity and can be used for gene therapy or in a vaccine. The nucleic acid, polypeptide encoded by it, or antibody to the polypeptide can be used in pharmaceutical compositions or vaccines for preventing or aborting pregnancy. PAPP-E is used in the antenatal diagnosis of dysgenetic pregnancies. The nucleic acids are used as probes to assess the level of PAPP-E isoform mRNA in chorionic villus samples, and the antibodies can be used to assess the expression levels of PAPP-E isoform proteins in chorionic villus samples, to diagnose dysgenetic pregnancies antenatally. This sequence represents an oligomer used in scanning the human PAPP-E genes described in the disclosure of the invention
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                                                                                                                                                                                                                                                                                                   New isolated nucleic acid encoding an isoform of human pregnancy associated plasma protein B, for preventing or aborting pregnanc
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contraceptive;
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(SHAN/) SHANNON M
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                                                                                                                                                                                                                                                                                                                             CC The present invention relates to novel human zinc finger-containing CC proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is CC encoded at chromosome 7q22.1, MDZ4 is encoded at chromosome 16p11.2 and MDZ12 is encoded at chromosome CC 15q26.1. The MDZ3, MDZ4, MDZ7, and MDZ12 sequences are useful in therapy, CC or in manufacturing a medicament for treating or preventing a disorder CC associated with decreased or increased expression or activity of MDZ3, CC MDZ4, MDZ7, or MDZ12, e.g. cancer or developmental disorders. The nucleic CC acids and proteins are also useful for diagnosing or monitoring a disease CC caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic CC acids can also be used as probes to detect and characterize gross CC electric in constructing microarrays for measuring gene expression. The CC proteins are useful as therapeutic agents for gene therapy or as CC vaccines. The present sequence was used to illustrate the invention.
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Matches 20; Conserv
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                                                                                                                                                                                                                                                                           Sequence 25 BP; 7 A; 7 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 8; SEQ ID NO 2988; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-AUG-2001; 2001US-00922181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human MDZ3 scanning oligonucleotide SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (AEOM-) AEOMICA INC.
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24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or MDZ12, e.g. cancer.
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                                                                                                                                         Conservative
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                                                                                                                                 Score 17.6; DB 1;
Pred. No. 4.5e+02;
0; Mismatches 4
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RESULT 316 ADB02001/c

Human microarray DNA oligonucleotide SEQ ID

14-OCT-2003

(first entry)

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RESULT 317
ACK21062/c
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Best Local S
Matches 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  proteins and their coding sequences: MDZ3, MDZ4, MDZ7, MDZ12. MDZ3 is encoded at chromosome 7g22.1, MDZ4 is encoded at chromosome 6g21.3-22.2, MDZ7 is encoded at chromosome 1fp11.2 and MDZ12 is encoded at chromosome 1fp11.2 and MDZ12 is encoded at chromosome 1fg11.2 and MDZ12 sequences are useful in therapy, or in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ3, MDZ1, or MDZ12, e.g. cancer or developmental disorders. The nucleic acids and proteins are also useful for disgnosing or monitoring a disease caused by altered expression of MDZ3, MDZ4, MDZ7, or MDZ12. The nucleic acids can also be used as probes to detect and characterize gross alterations in MDZ3, MDZ4, MDZ7, or MDZ12 genetic locus. The probes are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New zinc finger-containing proteins and nucleic acids, useful in manufacturing a medicament for treating or preventing a disorder associated with decreased or increased expression or activity of MDZ4, MDZ7 or MDZ12, e.g. cancer.
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                                                                                                                             ACK21062 standard;
                                                                                                                                                                                                                                                                                                                                                                            Sequence 25 BP; 7 A; 7 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                             useful in constructing microarrays for measuring gene expression. The proteins are useful as therapeutic agents for gene therapy or as vaccines. The present sequence was used to illustrate the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 8; SEQ ID NO 2987; 103pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shannon M, Gu Y, Nguyen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-JUL-2002; 2002EP-00016874.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to novel human zinc finger-containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-NOV-2003
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                                                                                                                             DNA;
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Pred. No. 4.
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RESULT 318
ACI40802/c
ID ACI408
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AC ACI408
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DT 13-OCT
XX
Human
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KW EST; [
KW genet;
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CR Also disclosed is a method of gene expression analysis. The array is used in monitoring gene expression levels by hybridisation to a DNA library, in analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis of at least one target sequence. The method of analysis comprises conclude acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises monitoring gene expression levels, identifying biallelic markers or polymorphisms, or family members of a gene and a cross-species comparison. Each of the nucleic acid further comprises at as sequence. The array of nucleic acid probes is useful in situ hybridisation, in Southern, Northern or dotblot hybridisation to identify or detect the sequence or specific mutations of any gene, in mapping the 5' termini of mRNA molecules by CC primer extensions or in screening cDNA or genomic libraries or subclones containing segments of DNA that have been concluded and previously sequence. The sequence of the isolated and previously sequence. The sequence of the collect acid probes incorporated in the microarray. Note: The sequence collect acid probes incorporated in the microarray. Note: The sequence collect acid probes incorporated in the microarray. Note: The sequence collect acid probes incorporated in the microarray of the collection of the collection of the microarray of the collection of the collection of the microarray of the collection of the collecti
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Best Local Similarity
  EST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New array of nucleic acid probes, useful for in situ hybridization, Southern, Northern or dot-blot hybridization to identify or detect t sequence or specific mutations of any gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its
                                                                           Human microarray DNA oligonucleotide SEQ ID NO 40793.
                                                                                                                                                                                                                       ACI40802 standard; DNA; 25 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 25 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AFFY-) AFFYMETRIX INC
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                                                                                                                                                                                                                                                                                                                                                                                    559 TTGGAGTTCCTGAAGAAGGAGGAG 582
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 17.6;
Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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CC perfect match, perfect mismatch, antisense match or antisense mismatch.

CC Also disclosed is a method of gene expression analysis. The array is used compounds. The nucleic acid probes are specifically designed for analysis of genetic variation or in hybridisation to a DNA library, compounds. The nucleic acid probes are specifically designed for analysis computed to a tleast one or more nucleic acids to at least two or more nucleic acids to at least two or more nucleic acids to at least two or more computed to a solid support. The analysis comprises monitoring gene expression levels, identifying biallalic markers or polymorphisms, cor family members of a gene and a cross-species comparison. Each of the culeic acids further comprises a tag sequence. The array of nucleic acid comparison seful in in situ hybridisation, in Southern, Northern or dotolot hybridisation to identify or detect the sequence or specific culeic acid probes is useful in in situ hybridisation, in Southern, Northern or dotolot hybridisations of any gene, in mapping the 5' termini of mRNA molecules by cor additional subclones containing segments of DNA that have been consolated and previously sequenced. The sequence presented is one of the culeic acid probes incorporated in the microarray. Note: The sequence containing the sequence presented is one of the solated and previously sequenced. The sequence for this patent can also be obtained in electronic format directly from USPTO at sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention discloses a microarray comprising a plurality of acid probes including one of 2,018,500 fully defined sequences,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 40793; 9pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New array of nucleic acid probes, useful for Southern, Northern or dot-blot hybridization sequence or specific mutations of any gene.
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to identify or detect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        or its
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Sequence 25 BP; 1 A; 7 C; 5 G; 12 T; 0 U; 0 Other;

Matches Query Match Best Local : 20; Similarity Conservative 83.3%; 0.3%; 0; Mismatches Score 17.6; Pred. No. 4. 4.5e+02 DB 1; Length 25; 0 Gaps 0

밁 8 RESULT 319 4399 AAAGACAAGAAAGATGAGACTCTG 4422

ACK27097

ACK27097 standard; DNA; 25

ВP

ACK27097

14-OCT-2003 microarray (first entry) DNA oligonucleotide SEQ

IJ

NO 127078.

EST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human;

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RESULT 320
ACK28333/c
ID ACK283
XX ACK283
AC ACK283
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DT 14-OCT
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DE Human
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EST; E
KW Geneti
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OS Homo E
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                                                                              EST; ss; probe; expressed sequence tag; microarray; gene expression; genetic variation; biallelic marker; polymorphism; human;
                                                                                                                                                                                                                                                                        ACK28333 standard; DNA; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for additional subclones containing segments of DNA that have been isolated and previously sequenced. The sequence presented is one of the nucleic acid probes incorporated in the microarray. Note: The sequence data for this patent can also be obtained in electronic format directly from USPTO at segdata.uspto.goc/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its perfect match, perfect mismatch, antiense match or antisense mismatch. Also disclosed is a method of gene expression analysis. The array is us
                                                                cross-species
                                                                                                                                            Human microarray DNA oligonucleotide SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence or specific mutations of any gene.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2003104410-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AFFY-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                 4547 GAGAGCAGCTGATAGCCCGAGTGA 4570
                                                                                                                                                                                                                                                                                                                                                                            AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                            GAGAGCAGCGGCTCGCCCGACTGA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                     (first entry)
                                                             comparison.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.3%;
                                                                                                                                                                                                                                                                        ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.6;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4.5e+02
                                                                                                                                               ID NO 128314.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         in situ hybridization, in
to identify or detect the
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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call probes including one of 2,018,500 fully defined sequences, or its perfect match, perfect mismatch, antisense match or antisense mismatch. Classification to a punk library, is used in monitoring gene expression levels by hybridisation to a DNA library, is used compounds. The nucleic acid probes are specifically designed for analysis of genetic variation or in hybridisation of tag-labelled compounds. The nucleic acid probes are specifically designed for analysis comprises comprises acid probes and detecting the hybridisation. The nucleic acid probes and detecting the hybridisation. The nucleic acid probes are attached to a solid support. The analysis comprises mucleic acid grobes are attached to a solid support. The analysis comprises monitoring composes are attached to a solid support. The analysis comprises monitoring confamily members of a gene and a cross-species comparison. Each of the nucleic acids further comprises a stag sequence. The array of nucleic acid probes is useful in in situ hybridisation, in Southern, Northern or dothot hybridisation to identify or detect the sequence or specific cutations of any gene, in mapping the 5' termini of mRNA molecules by primer extensions or in screening cDNA or genomic libraries or subclones for additional subclones containing segments of DNA that have been isolated and previously sequenced. The sequence presented is one of the full for this patent can also be obtained in electronic format directly from USPTO at segdata.uspto.goc/sequence.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 128314; 9pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New array of nucleic acid probes, useful for Southern, Northern or dot-blot hybridization sequence or specific mutations of any gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-MAR-2002; 2002US-00098263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2003104410-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention discloses a microarray comprising a plurality of nucleic acid probes including one of 2,018,500 fully defined sequences, or its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-MAR-2001; 2001US-0276759P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AFFY-) AFFYMETRIX INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2003-567953/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            in situ hybridization,
to identify or detect
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Query Match Best Local S Matches 20 Local Similarity 0.3%; ilarity 83.3%; Conservative BP; 3 A; 9 C; 8 G; 5 T; 0 U; 0 Other; <u>,</u> Score 17.6; DB 1; Pred. No. 4.5e+02; Mismatches Length 25;

0

Gaps

0

Sequence 25

밁 Ś 4547 GAGAGCAGCTGATAGCCCGAGTGA 4570 25 GAGAGCAGCGGCTCGCCCGACTGA 2

ACH54326 RESULT 321 ACH54326; ACH54326 standard; DNA; 25 (first entry)

16-OCT-2003

DNA target sequence #3462 useful in array for genetic analyses.

Gene expression analysis; array; hybridisation; genetic variation; tag-labelled compound; gene family; in situ hybridisation; library screening; Southern hybridisation; northern hybridisation; dot-blot hybridisation; gene sequence; mutation detection; target sequence; probe; PCR; primer; ss.

Unidentified

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RESULT 322
ADK65715/c
ID ADK657
XX
ADK657
XX
AC ADK657
XX
VX
AC HAAP
XX
DE A thal
XX
Vitami
KW food p
XX
Food p
XX
OS Arabid
XX
DE1021
XX
DE1021
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                         8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to nucleic acid sequences that are complementary to particular genes, and can be used as probes for a complementary to particular genes, and can be used as probes for a comprises 9 or more consecutive nucleotides from at least one of 14936 nucleotide sequences defined in the patent, or their perfect sense match, sense mismatch, antisense match or antisense mismatch oligonucleotides. The probes may be used in an array comprising at least 10 distinct nucleic acid probes. The array is useful in monitoring gene expression levels by hybridisation to a DNA library, in analysing genetic variations, and in hybridisations to a blood libraries of a gene. The probes are useful for identifying family members of a gene. The probes are also useful in in situ hybridisations, in screening cDNA or genomic libraries (or derived subclones) for additional clones containing segments of DNA that have been previously isolated and sequenced, in Southern, northern, or dot-blot hybridisation of genomic DNA to identify or detect the sequence of any gene or detect specific mutations in any gene, and in mapping the 5' termin of mana molecules by primer extensions. The nucleic acid sequences of the invention are also useful as PCR primers. The invention provides a large collection of nucleic acid sequences concerns the target sequences of the invention. Note: The sequence data for this patent was obtained in electronic format the library than a sequence of the invention. Note:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 20
                                                                                                                    vitamin E; 2-methyl-6-phytylhydroquinone methyltransfase; enzyme; food processing; abiotic stress; ss; primer; PCR.
                                                                                                                                                                                                                       06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New probe array useful e.g. for monitoring gene expression levels, for analyzing genetic variations, or for hybridizing tag-labeled compounds, comprises multiple nucleic acid probes.
                                                                              Arabidopsis thaliana
                                                                                                                                                                              A thaliana 2-methyl-6-phytylhydroguinone methyltransfase PCR primer
                                                                                                                                                                                                                                                             ADK65715;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 25 BP; 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim
                                         DE10212703-A1
                                                                                                                                                                                                                                                                                                    ADK65715 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-576608/54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-AUG-2001; 2001US-0311040P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-AUG-2002; 2002US-00215112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MITT/) MITTMANN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US2003082596-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                           4293 CTCCATTCGGAAGGAACTGGAGCT 4316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1, SEQ ID NO 3462; 9pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      CTCCTTACGGAAGGAACAGGACCT 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 7 C; 6 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USPTO web site at seqdata.uspto.gov/psipsDIDEntry.html
                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 17.6; DB 1;
Pred. No. 4.5e+02;
0; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Length 25;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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RESULT 323
ADO81067/c
ID ADO810
XX Gene t
XX Gene t
XX Gene t
XX Cystic
XX Milk p
XX Milk p
XX Milk p
XX Milk p
XX Gene t
XX G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  문
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a method for preparing Vitamin E, which comprises culturing an organism that has higher 2-methyl-6-phytylhydroquinone methyltransferase activity than the wild type. The method is used to produce transgenic organisms, particularly plants that have increased Vitamin E content, useful in foods, fodder (or supplements), for food processing, for preparing Vitamin E-containing extracts, and for converting 2-methyl-6-(phytyl, solanesyl or geranylgeranyl) hydroquinones to the corresponding 2,3-dimethyl compounds. Also, the method is used for producing plants with increased Vitamin E content which have better resistance to abiotic stress. The present sequence is a PCR primer used to isolate the A. thaliana 2-methyl-6-phytylburger.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic dimilk protein; hormone; transcription factor; pT7-blue-vector;
                                  WPI; 2004-215730/21
                                                                                          Geldermann H,
                                                                                                                                                                                                             09-AUG-2002; 2002DE-01036711.
                                                                                                                                                                                                                                                                     09-AUG-2002; 2002DE-01036711.
                                                                                                                                                                                                                                                                                                                                26-FEB-2004.
                                                                                                                                                                                                                                                                                                                                                                                           DE10236711-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 microsatellite; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene typing; polymorphic microsatellite loci; PML;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cow prion protein microsatellite locus primer #79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADO81067;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADO81067 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 25 BP; 4 A; 9 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             phytylhydroquinone-methyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 35; 48pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   phytylhydroquinone methyltransferase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Preparation of Vitamin E, useful e.g. in foods and animal fodder, comprises growing organisms, with increased activity of 2-methyl-6-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-215445/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-MAR-2002; 2002DE-01012703
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (SUNG-) SUNGENE GMBH & CO KGAA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1201 TCAGAGAAAGGCAGGCCCCCATGG 1224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20;
                                                                                                                                                     UNIV HOHENHEIM.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGAGAGTAAGGTCGGCCCCCATGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                          Preuss S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              83.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Salchert K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25
                                                                                             Han
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 17.6;
Pred. No. 4.
                                                                                          ĸ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Geiger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4.5e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
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0,

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ARESULT 324
AAZ71491/c
ID AAZ714
XX AAZ714
XX AAZ714
XX Human
XX Human
XX Human
XX Human
XX Homo s
XX Claim
XX Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         more PML; and prediagnosis (M3) of diseases associated with gene that include PML. The method is used to identify microsatellite markers, in a disease-related gene, that are associated with a predisposition to diseases and for prediagnosis of such diseases, especially prion diseases but also cystic fibrosis, malignant hyperthermia syndrome in pigs and metabolic diseases; also to type genes that encode milk proteins, hormones or transcription factors. The method is simpler, quicker and particularly less expensive than known methods based on sequencing. This sequence represents a primer used to genotype a region of the cow prion protein (PrP) comprising a polymorphic microsatellite locus.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide nolymorphic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention describes a method of typing (M1) a gene (I) that has one or more polymorphic microsatellite loci (PML). The method comprises PCR amplification of at least one DNA region of (I) that includes PML, using as template a DNA sample containing at least one segment of (I); and determining the length of the resulting amplicon(s). Also described are: a method of determining (M2) microsatellite markers (MM) for predisposition to a disease, associated with a gene that includes one or
                                                                                                                                                                                                                      21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human biallelic marker upstream amplification primer SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                           Novel biallelic markers used
                                                                                                              WPI; 2000-013267/01.
                                                                                                                                              Cohen D,
                                                                                                                                                                                                                                                                                                                28-OCT-1999
                                                                                                                                                                                                                                                                                                                                                  WO9954500-A2
                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                       diagnosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ71491;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ71491 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; Page 28; 64pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Typing genes that contain polymorphic microsatellite loci, useful for identifying predisposition to disease, by amplification and determining length of amplicons.
                                                                                                                                                                                    (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                          of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5398 AATACAAAAAAGAAAAATGAAAA 5421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAAAAAAAAAGAAAGAAAGAAAA 2
                                                                                                                                              Blumenfeld M,
                                                        human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                    98US-0082614P
98US-0109732P
                                                                                                                                                                                                                                                                            99WO-IB000822
                                                        genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
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                                                                                                                                                Chumakov
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This invention describes a novel oligonucleoside compound for hybridising to a target nucleic acid strand. The oligonucleoside comprises (a) an oligonucleoside sequence that is complementary to a target region or subregion of the target nucleic acid strand and (b) a portion that is non complementary to a target site in the target region or subregion such that, when the oligonucleoside compound is hybridised to the target strand, a base group at the site is oriented away from an inter-strand orientation. The oligonucleoside and combinations are used for inhibiting production of a selected protein in a cell by effecting cleavage at a site in a target region of cellular RNA that codes for the selected protein. They can be used for treating a condition in a mammal that is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AA265654 to AA269578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AA269579 to AA277440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not accually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                            Oligo:nucleoside compounds for cleaving RNA - complementary to a target nucleic acid strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cleavage; oligonucleoside; target; inter-strand orientation; inhibitor; disease; treatment; internucleotidyl bond cleavage; primer; ss.
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                                                                                                                                                                                                                       Disclosure; Page 92; 109pp; English
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Pred. No. 4.
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The invention relates to an isolated polypeptide, comprising a variant form of mouse or human SAC1 polypeptide. The variant form is associated with altered preference for carbohydrates, other sweeteners or ethanol. The polypeptide and its associated DNA sequence can be produced by recombinant techniques and is useful for preventing obesity, diabetes or alcoholism associated with SAC1 expression. The sequences are useful in screening for drugs and sweeteners. Recombinant cell lines and transgenic embryos may be used in screening for and identifying agents that induce or repress function of SAC1. Predisposition to diabetes, obesity or alcoholism can be ascretained by testing any fluid or tissue of a human (such as blood, pancreas or tongue) for sequence variations of the SAC1 gredisposition to diabetes, obesity and/or alcoholism and may provide a diagnostic mark. The polymucleotide can be detected in a biological sample by contacting the DNA with a probe to form a hybridisation complex
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Ohmen JD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 14; Page 89; 239pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel isolated polypeptide comprising variant form of mouse or human SACI polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.
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28-JUL-2000; 2000US-0221419P.
10-NOV-2000; 2000US-0247443P.
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94.7%;
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Tordoff MG;
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                            The invention relates to an isolated polypeptide, comprising a variant form of mouse or human SAC1 polypeptide. The variant form is associated with altered preference for carbohydrates, other sweeteners or ethanol. The polypeptide and its associated DNA sequence can be produced by recombinant techniques and is useful for preventing obesity, diabetes or alcoholism associated with SAC1 expression. The sequences are useful in screening for drugs and sweeteners. Recombinant cell lines and transgenic embryos may be used in screening for and identifying agents that induce or repress function of SAC1. Predisposition to diabetes, obesity or alcoholism can be ascertained by testing any fluid or tissue of a human (such as blood, pancreas or tongue) for sequence variations of the SAC1 (such as blood, pancreas or tongue) for sequence variations of the SAC1 (such as blood, pancreas or tongue) for sequence variations of the SAC1 (such as blood, pancreas or tongue) for sequence variations of the SAC1 (such as blood, pancreas or tongue) for sequence variations of the SAC1 (such as blood, pancreas or tongue) for sequence variations of the SAC1 (such as blood, pancreas or tongue) for sequence variations of the SAC1 (such as blood, pancreas or tongue) for sequence variations of the SAC1 (such as blood) for the SAC1 (su
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Ohmen JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel isolated polypeptide comprising variant form of mouse or human SAC: polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-APR-2000; 2000US-0200794P
28-JUL-2000; 2000US-0221419P
10-NOV-2000; 2000US-0247443P
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blood; tongue; PCR primer; anorectic; antidiabetic; gene therapy;
protein replacement therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Reed
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene-specific oligonucleotide PCR primer #419
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1 DR, Ross D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LAMBERT CO.
CHEM SENSES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         carbohydrate; sweetener; ethanol; alcoholism;
transgenic embryo; body tissue; body fluid; par
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Pred. No. 4.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chatterjee A,
Tordoff MG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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RESULT 328
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                                                                                                                                                                                   cc insulin-like growth factor binding protein-3 (IGRPP-3) interacting a competition, termed P4.33 protein, IGRPP-3 is used in gene therapy. Antibody cc specific for P4.33 is useful in an assay for cancer treatment, prognosis, cc or diagnosis in a patient for cancer cells that express P4.33 protein or peptides in a patient for cancer cells that express P4.33 protein or peptides in competides, by determining the amount of P4.33 protein or peptides in cc blood, serum, urine, lymph, saliva, tumour tissue, placental tissue, umbilical cord tissue, amnioric fluid, chorionic vill tissue or their combinations, by enzyme linked immunosorbent assay (ELISA), Western canalysis, immunoprecipitation or immunohistocytochemistry. Detecting P4.33 specific sequences in a bodily fluid sample from a patient is also cuseful in an assay for cancer treatment, prognosis, or diagnosis in a competition of cancer cells that express P4.33-specific sequences, by performing a sequence identity assay such as ELISA immunologic assays, cordinated prognosis, or diagnosis in a coperific sequences. P4.33 is useful for preventing or treating cancer, concluding lung, cervical, breast, colon or prostate carcinoma in a coperific sequences. P4.33 is useful for preventing or treating cancer, conhibition of DNA synthesis and cellular proliferation and in the cinduction of apoptosis in cancer cells. The present sequence is human conduction of apoptosis in cancer cells.
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel insulin-like growth factor binding protein-3 interacting protein, termed P4.33 for identifying compounds having anti-cancer activity, inducing apoptosis and inhibiting cellular proliferation in cancer cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; insulin-like growth factor binding protein-3; IGFBP-3; cytostatic; lung; cervical; breast; colon; cancer; prostate carcinoma; P4.33 protein; gene therapy; cellular proliferation; receptor; apoptosis; antisense; ss.
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                                                                                                    Sequence 20 BP; 1 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 47; Page 18; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-MAY-2001; 2001WO-US016437.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to an isolated insulin-like growth factor binding protein-3
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0.3%;
                                                                                                13 C; 2 G; 4 T; 0 U; 0 Other;
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Score 17.4; DB 1; Length 20; Pred. No. 4.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ONA sequence encoding a (IGFBP-3) interacting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
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RESULT 329
ABS74301
ID ABS743
XX OP-DEC
XX Human;
XW Gene t
XW Worsin-
KW Huntin
KW Mypoxii
CX Thomo s
XX US6441
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                                                                                     The invention relates to a purified nucleic acid sequence (referred as CC CACNA2D2 gene which encodes human calcium channel alpha2delta-2 subunit CC protein) comprising a fully defined alpha2delta splice isoform 1, 2 or 3 cc nucleic acid sequence, or its complement and the encoded proteins. Also include are: (1) a method of producing a calcium channel protein which comprises introducing a recombinant expression vector comprising the calcium channel protein, into a cultured host cell under conditions such that the host cell expresses the amino acid sequences; and (2) a method for co-expressing calcium channel proteins, comprising carrying out the method of (1), but with one or more canced that one expression vector comprising one or more nucleic acid sequences canceding the splice variants. CACNA2D2 nucleic acid is useful for correct than one expression vector comprising one or more nucleic acid sequences canceding the splice variants. CACNA2D2 nucleic acid is useful for comprising one or more nucleic acid sequences conjupeptide is useful for treating patients with Lambert-Saton myasthenic syndrome (LEMS) (an autoimmune disease) and for identifying compounds cuseful for treating other diseases associated with abnormal calcium channel protein activity (e.g. epilepsy, migraine, episodic ataxia, cardiac arrhythmia. angina
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel purified nucleic acid sequence encoding human calcium channel alpha2delta subunit protein, useful for detecting, preventing and treating cancer, stroke, brain trauma, Huntington's disease, myocardial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lerman MI, Latif F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-DEC-1998;
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herapy; Lambert-Eaton myasthenic syndrome; LEMS; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 47; 77pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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seizures, Huntington's disease, amnesia, cardiac arrhythmia, angina pectoris, hypoxic damage to the cardiovascular system, ischaemic dar to the cardiovascular system, ischaemic dar

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RESULT 330
ACD25699
ID ACD256
XX ACD256
XX ACD256
XX ACD256
XX Human;
XW Human;
XW Human;
XW Carvic
XW Darin
XW Amnesi
XW Single
XX US2003
XX O6-MAN
XX O6-MAN
XX (LERM;
PA (LATI)
PA (LATI)
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                       (LERM/) LERMAN M I.
(LATIF F.
(WEIM/) WEI M.
(DUHF/) DUH F.
(MINN/) MINNA J D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; 88; PCR; calcium channel alpha2delta; chromosome 3p21.3; primer; transgenic; cancer; lung cancer; small cell carcinoma; epilepsy; stroke non-small cell carcinoma; breast cancer; nasopharyngeal cancer; cervical cancer; head and neck cancer; neurological disease; brain trauma; Alzheimer's disease; multiinfarct dementia; seizure; amyotrophic lateral sclerosis; convulsions; Huntington's disease; amyotrophic cardiovascular disease; cardiac arrhythmia; angina pectoris; hypoxic damage; ischaemia; myocardial infarction; SSCP; congestive heart failure; Lambert-Earon myasthenic syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              failure, muscular dystrophy and hypertension) CACNA2D2 nucleic acid is useful as primers and probes for detecting presence of nucleic acid sequence encoding at least a portion of calcium channel protein, in detection, identification and isolation of alpha2delta sequences diagnosing and typing of preneoplasias and cancers, since genetic disruption of 3p21.3 region (in which the alpha 2delta gene is located) is common in cancer (e.g. lung cancer and breast cancer) and preneoplastic lesion (e.g. hyperplasia, dysplasia, carcinoma in situ). The present is an SSCP (single strand change polymorphism) PCR primer used to detect polymorphisms in sequences encoding a human calcium channel alpha2delta splice isoform protein
                                                                                       New substantially pure human calcium channel alpha2delta subunit splice isoform 1, 2 and 3 sequence useful in preventing, treating and diagnosis cancer, neurological disorders and cardiovascular disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-DEC-1998;
22-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human calcium channel alpha2delta SSCP primer MJE8F.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACD25699 standard;
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                                                                                                                                                                                                                                                                          Lerman MI,
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                                                                                                                                                                                                                                                                                                                                     SEKIDO Y.
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                                                                                                                                                                                                                                                                          Latif F,
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99US-00470443.
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94.7%;
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Pred. No. 4.6e+02;
                                                                                                                                                                                                                                                                          Duh P,
                                                                                                                                                                                                                                                                          Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 20
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Example 7;

Page 25;

79pp; English

Novel single or multiple target oligonucleotide anti-sense to e.g.

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AC ADJ613
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC complishing at least a portion of human calcium channel alpha2delta control splice isoform 1, splice isoform 2 sequence or splice isoform 3, cor their variants, and their encoding nucleic acids (or their complements, variants, or homologues). Also included are screening a test compound for modulating calcium channel activity, an antibody which binds to the calcium channel or its variants and producing a transgenic non-channel alpha 2delta subunit relative to a corresponding wild-type canimal. The calcium channel proteins are useful for generating an antibody (which is useful for detecting the proteins or their portions). The transgenic animal (preferably a rodent e.g. mouse) is useful for identifying a therapeutic compound for treating a transgenic animal carcinoma or non-small carcinoma). The daracting a neurological disease, especially epilepsy, campotrophic lateral sclerosis, convulsions, seizures, Huntington's carcive hear a cardiovascular disease, especially cardiac arrhythmia, angina pectoris, hypoxic damage to the cardiovascular system, myocardial infarction, and congestive heart failure; or Lambert-Saton myasthenic syndrome. The proteins and nucleic acids are useful in the diagnosis, prevention and conformation polymorphism) primer used to detect colymorphisms in the calcium channel alpha2delta subunit gene
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           interleukin; II-4 receptor; II-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide associated to IL5R-X61176 #15
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                                                          WPI; 2004-203534/19
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                                                                                                                                                                                                        (EPIG-) EPIGENESIS PHARM INC.
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lin S,
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                                                                                                               Ę,
                                                                                                               Sandrasagra A,
H, Cong H;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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                                                                                                                                           Aguilar
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RESULT 332
ADJ61322/c
ID ADJ613
XX ADJ613
XX ADJ613
XX Interl
KW interl
KW interl
KW sirway
KW cystic
KW pulmon
KW pulmon
KW pulmon
KW ES-JUL
XX ADJ613
XX INTERL
XX I
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Best Local Similarity
Matches 18; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene(e) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is useful for production of a medicament for the prevention and/or treatment of a respiratory or lung disease. The respiratory or lung disease is chosen from airway inflammation, allergy(ies), asthma, impeded respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                          Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant gene CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide associated to IL5R-X61176
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                                                                                                                                                                                                                                                                                                                           EPIG-)
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                                                      e.g.,
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in s,
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H, Cong H;
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Pred.
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No. 4.
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                                                                                                           genes e.
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NO 2178;

85pp;

The present invention relates to an oligonucleotide anti-sense to eninitiation codon, coding region with 2-10 nucleotides of 5'-end and end of nucleic acid target comprising gene(s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the

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RESULT 333
ADJ61324/c
ID ADJ613
XX ADJ613
XX ADJ613
XX ADJ613
XX Oligon
XX interl
KW airway
KW cystic
KW pulmon
KW ss.
XX Homo s
XX Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cc initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-
cc initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-
cc end of nucleic acid target comprising gene(s) chosen from e.g.
cc interleukin (II)-4 receptor, II-5 receptor or salts of the
cc oligonucleotide and optionally surfactant operatively linked to the
cc oligonucleotide. The method is useful for preventing or treating a
cc respiratory or lung disease, which involves administering to the airways
cc of a subject an effective amount of an inhibitor. The oligonucleotide is
cc useful for production of a medicament for the prevention and/or treatment
cc of a respiratory or lung disease. The respiratory or lung disease is
cc chosen from airway inflammation, allergy(ies), asthma, impeded
cc respiration, cystic fibrosis (CF), chronic obstructive pulmonary diseases
cc (OOPD), allergic rhinitis (AR), acute respiratory distress syndrome
cc (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway
cc obstruction. The present sequence represents an oligonucleotide of the
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Best Local S
Matches 18
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airway inflammation; allergy; asthma; impeded respiration;
cystic fibrosis; acute respiratory distress syndrome;
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                                                                                                                             Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide associated to IL5R-X61176 #16.
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                                                                                            Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pulmonary hypertension; lung inflammation;
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                                                                                                                                                                                                                                                           Shahabuddin
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in S,
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                                                                                            ID NO
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                                                                                                                                                                                                                                                           Sandrasagra
H, Cong H;
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Pred. No. 4.6e
0; Mismatches
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                        The present invention relates to an oligonucleotide anti-sense to e.g., initiation codon, coding region with 2-10 nucleotides of 5'-end and 3'-end of nucleic acid target comprising gene (s) chosen from e.g. interleukin (IL)-4 receptor, IL-5 receptor or salts of the oligonucleotide and optionally surfactant operatively linked to the oligonucleotide. The method is useful for preventing or treating a respiratory or lung disease, which involves administering to the airways of a subject an effective amount of an inhibitor. The oligonucleotide is
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                                                                                                                                                                                                                                                                                                                                           Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
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                                                                                                                                                                                                                                                                   ID NO 2181; 85pp; English
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RESULT 336 ADO46712/c ID ADO467

ADO46712 standard; DNA;

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RESULT 335
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                                                                                                                                                                                                                                                The present invention relates to identifying a rice variety as amplification genetic marker and identifying whether test rice variety one of the 32 rice varieties e.g., Kasalath, breath which can Hayamasari, Italica Livorno, Dunghan Shali, Arroz Da Terra, Fany Mihombare. The method is useful for identifying rice variety and identifies excellent rice variety. The present sequence represent containing SSR sequence of the invention.
                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amplifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2004-003560/01
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 22;
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HOKKAIDO GREEN BIO KENKYUSH
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genetic marker.
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                                                                                                   Conservative
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Pred. No. 4.
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CC codon, coding region, 5' or 3' intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (IL)-4 receptor; interleukin (IL)-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 CO repE4 D. The invention CC also relates to a method of screening a candidate compound that binds to come or more nucleic acid target(s) or expressed product(s), for the CC one or more nucleic acid target(s) or inhibiting expression of a conjugate or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are cusful for preventing or treating a respiratory or lung disease. The CC tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are CC tryptase b, and/or sathma and/or lung disease. The CC respiratory or lung disease is associated with hyper-responsiveness to captor increased levels of, adenosine and/or levels of adenosine A CC receptor(s), and/or sathma and/or lung allergies associated with CC inflammation or an inflammatory disease. The respiratory or lung disease (CDD), chronic obstructive pulmonary disease (CDD), chronic obstructive pulmonary disease (CDD), chronic obstructive pulmonary disease of the bronchocreativities of the component of the comp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to codon, coding region, 5'
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23-APR-2002;
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pronchoconstriction.
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MILLER S.
SHAHABUDDIN
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CONG H.
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TANG L.
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24, useful for
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This sequence represents an oligonucleotide
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of the
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RESULT 337
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Best Local S
Matches 18
chosen from a gene encoding interleukin (IL)-4 receptor, interleukin (IL)-5 receptor, CCR1, CCR3, Ectaxin-1, RAMTES, MCP4, CD23, ICAM, VCAM, tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention also relates to a method of screening a candidate compound that binds to one or more nucleic acid target(s) or expressed product(s), for the prevention and/or treatment of a respiratory or lung disease. The
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                                                                                                                                                                 The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target
                                                                                                                                                                                                                                                                                  Claim 2;
                                                                                                                                                                                                                                                                                                                                                                Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
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23-APR-2002; 2002WO-US013143.
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CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a;
tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease;
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(TANG/)
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MILLER S.
SHAHABUDDIN S.
LU H.
CONG H.
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TANG L.
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                                                                                                                                                                                                                                                                                  SEQ ID NO 2182; 174pp; English
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in S. Lu H, Cong
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ong H;
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Pred. No. 4.
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oligonucleotides are useful for reducing or inhibiting expression of a

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RESULT 338
AD046715/c
ID AD0467
XX AD0467
XX AD0467
XX AD0467
XX Human;
XW Human;
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Matches 18
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23-APR-2002; 2002WO-US013143.
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Shahabuddin S, Lu H, Cong
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                                                                           WPI; 2004-293804/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammation; bronchitis; airway obstruction; bronchoconstriction
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                                                                                                                                                                                                                                                          CONG H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NYCE J W.
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                                                                                                                                                                                                                                                                                                                                  SHAHABUDDIN S.
                                                                                                                                                                                                                                                                                                                                                                        AGUILAR D.
MILLER S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TANG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SANDRASAGRA A.
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ng H;
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                                                                                                                                                                                  Aguilar D,
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initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
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Claim SEQ ID NO 2181; 174pp; English.

CC chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II)CC chosen from a gene encoding interleukin (II)CC tryptage a, tryptage b, PD84 A, PD84 C or PD84 D. The invention
CC tryptage a, tryptage b, PD84 A, PD84 C or PD84 C or PD84 D. The invention
CC also relates to a method of screening a candidate compound that binds to
CC one or more nucleic acid target(s) or expressed product(s), for the
CC prevention and/or treatment of a respiratory or lung disease. The
CC prevention and/or treatment of a respiratory or lung disease in the CC oligonuclectides are useful for reducing or inhibiting expression of a
CC cral, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, tryptage a,
CC cryptage b, PD84 B, PD84 C, or PD84 D. The oligonuclectides are
CC useful for preventing or treating a respiratory or lung disease. The
CC respiratory or lung disease is associated with hyper-responsiveness to
CC and/or increased levels of, adenosine and/or levels of adenosine A
CC receptor(s), and/or asthma and/or lung allergies associated with
CC is chosen from airway inflammatory disease. The respiratory or lung disease
CC is chosen from airway inflammatory disease. The respiratory or lung disease
CC systic fibrosis (CF), chronic obstructive pulmonary disease (Opp),
CC hypertension, lung inflammation, bronchitis, airway obstruction or the codon, coding region, 5' or 3' intron-exon junction, intron or region with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target The invention relates to oligonucleotides anti-sense to an initiation codon, coding region, 5' or 3' intron-exon junction, intron or region bronchoconstriction. This sequence represents an oligonucleotide

Sequence 20 BP; 0 A; 10 C; 0 G; 10 T; 0 U; 0 Other;

S Query Match Best Local S Matches 18; Similarity Conservative 0.3%; 0; Mismatches Score 17.4; Pred. No. 4. 6e+02 DB 1; Length 20 Indels 0 0

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15-JUL-2004 (first entry) ADO46713 standard; DNA; 20 ВP

Human oligonucleotide #2079.

ADG46713/c
ID ADG46713/c
ID ADG467
XX ADG467
XX ADG467
XX ADG467
XX ADG467
XX Human
XX Human
XX Human
XX Human
XX CCR1;
XW CCR1;
XW CCR1;
XW CCR1;
XW CCR1;
XW Lrypta
XW Asrima
XW Asrima
XW Asrima
XW Iung 1
XX Chroni
XW Acute
XX US2004
XX US2004
XX US2004
XX US2004
XX US2004
XX STULL
XX 23-APR
PF 23-APR
PR 23-APR
XX 23-APR Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-CR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDB4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; chronic obstructive pulmonary disease; COPD; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; acute respiratory distress syndrome; pulmonary hypertension; 25-JUL-2003; 2003US-00627930. Homo sapiens US2004049022-A1 inflammation; bronchitis; airway ÇÊ

Novel single or multiple target oligonucleotide anti-sense to e.g.

23-APR-2002; 2002WO-US013135. 23-APR-2002; 2002WO-US013143.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to oligonucleotides anti-sense to an initiation CC codon, coding region, 5' or 3' intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (II) 4 receptor, interleukin (II).

CC tryptase a, tryptase b, PDE4 A, PDE4 B, PDE4 C or PDE4 D. The invention CC also relates to a method of screening a candidate compound that binds to CC one or more nucleic acid target(s) or expressed product(s), for the CC or or or more nucleic acid target(s) or expressed product(s), for the CC or or or many encoding interleukin-4 receptor, initiating expression of a CC gene or mRNA encoding interleukin-4 receptor, interleukin-5 receptor, CCR1, CCR3, Ectaxin-1, RANYES, MCP4, CD23, ICAM, VCAM, tryptase a, CC tryptase b, PDE4 A, PDE4 B, PDE4 C, or PDE4 D. The oligonucleotides are useful for preventing or treating a respiratory or lung disease. The CC espiratory or lung disease is associated with hyper-responsiveness to CC end/or increased levels of, adenosine and/or levels of adenosine and/or levels of adenosine and/or levels of adenosine and/or lavels of schosen from airway inflammation, allergy, asthma, impeded respiration, CC is chosen from airway inflammation, allergy, asthma, impeded respiration, CC inflammation, lavels by a struction or an inflammation, bronchitis, airway obstruction or the bronchitis, acute respiratory distress syndrome, pulmonary inflammation, bronchitis, airway obstruction or the bronchitis acute respiratory distress syndrome, pulmonary in the complex of the complex of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                       Human; ss; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease;
asthma; lung allergy; inflammation; inflammatory d: airway inflammation; allergy; impeded respiration;
                                                                                                                                                                                                                                                                         ADO46714 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
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                                                                                                                                                 Human oligonucleotide #2080
                                                                                                                                                                                            15-JUL-2004
                                                                                                                                                                                                                                     AD046714;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bronchoconstriction. This sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 2; SEQ ID NO 2179; 174pp; English.
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(SAND/)
(TANG/)
(AGUI/)
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(SHAH/)
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CONG H.
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SANDRASAGRA A.
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in S, Lu H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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Cong,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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Pred. No. 4.6e+02;
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      cystic fibrosis; CF,
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                                                                              Query Match
Best Local S
Matches 18
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                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codon, intron of respiratory disease-relevant gene e.g. CCR1, RAWTES, MCP4, useful for prophylaxis or treating respiratory disease e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                      allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction obsoronchoconstriction. This sequence represents an oligonucleotide observations.
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23-APR-2002; 2002WO-US013143.
                                                                                                                                                                                                     invention.
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TANG L.
AGUILAR D.
                                                                                                 Similarity
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                                        AGAGAAAGAGAGAGAGA 1198
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in S, Lu H, Cong
                                                                                                                                                            BP; 0 A;
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)ng H;
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                                                                              Pred. No. 4.6
                                                                                                 Score 17.4;
Pred. No. 4.
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                                                                                es 1;
                                                                                                                      DB 1; Length 20;
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                                                                                                                                                                 Other;
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RESULT

ADM16170 standard;

DNA;

20 BP

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ADM16170/c
ID ADM161
AC MOUSE;
KW MO
                                                                                      CC relates to a method for analysing a biomolecule in a biological sample, comprising altering SAC1 activity in the sample and measuring the activity, a method for analysing a polynucleotide in a biological sample, comprising contacting a polynucleotide in a biological sample, comprising contacting a polynucleotide in a biological sample with a grobe where the probe hybridises to a SAC1 polynucleotide to form a complex and detecting the hybridisation complex, a method of identifying susceptibility to obesity or diabetes comprising comparing the nucleotide sequence of the suspected SAC1 allele with a wild type concleotide sequence, where the difference between the suspected allele cand the wild-type sequence identifies a sequence variation of the SAC1 nucleotide sequence, and a method of treating or preventing obesity, comparising compositions or alcoholism associated with expression of SAC1, comprising composition and a transgenic administering to a subject a pharmaceutical composition and a transgenic cof the invention are useful for screening drugs for inhibition or crestoration of gene function as antidiabetic, antiobesity or antialcohol consumption therapies and for identifying sweeteners and alcohols. This sequence represents a PCR primer used to amplify murine SAC1 DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated polynucleotides for sensing carbohydrates, other sweeteners, or ethanol, useful for screening drugs for inhibition or restoration of gene function as antidiabetic, antiobesity or antialcohol consumption
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to SAC1 polypeptides and the polynucleotides encoding them. The polynucleotides contain a variation associated with sensing carbohydrates, other sweeteners or ethanol. The invention also
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 12; SEQ ID NO 440; 148pp; English
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ROSS D A.
OHMAN J D.
CHATTERJEE A.
DE JONG P J.
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BEAUCHAMP G
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De Jong PJ;
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Query Match Best Local Similarity

0.3%; 94.7%;

Score 17.4; DB 1; Pred. No. 4.6e+02;

Length

Sequence

20

BP; 4 A; 5 C; 6 G; 5 T; 0 U; 0 Other;

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RESULT 342
ADM16192/c
IDM ADM1611
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ADM1611
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ADM1611
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ADM161
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ADM16
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ADM1
The invention relates to SAC1 polypeptides and the polynucleotides concoding them. The polynucleotides contain a variation associated with concoding carbohydrates, other sweeteners or ethanol. The invention also corelates to a method for analysing a biomolecule in a biological sample, comprising altering SAC1 activity in the sample and measuring the cativity, a method for analysing a polynucleotide in a biological sample, comprising contacting a polynucleotide in a biological sample, comprising contacting a polynucleotide in a biological sample, comprising contacting a polynucleotide in a biological sample, comprising complex and detecting the hybridisation complex, a method of identifying susceptibility to obesity or diabetes comprising comparing the nucleotide sequence of the suspected SAC1 allele with a wild type can discotide sequence, where the difference between the suspected allele nucleotide sequence identifies a sequence variation of the SAC1 nucleotide sequence and a method of treating or preventing obesity, candinistering to a subject a pharmaceutical composition and a transgenic animal that carries an altered SAC1 allele. The methods and compositions of the invention are useful for screening drugs for inhibition or antialcohol crestoration of gene function as antidiabetic, antiobesity or antialcohol
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CHATTERJEE A.
DE JONG P J.
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BEAUCHAMP G
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an JD, Chatterjee
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antidiabetic; alcohol; anorectic; antialcoholic;
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PJ;
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RESULT 343
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Best Local Similarity
Matches 18; Conserv
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aziridinylcytosine; cross-linking group; tumour necrosis factor;
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/mod_base= OTHER
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                                         "N4N4-ethanocytosine"
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Pred. No. 4.6e
0; Mismatches
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RBSULT 344
AAQ20034/c
ID AAQ20034/standard; DNA; 21 BP
XX
AC AAQ20034;
XX
DT 01-APR-1992 (first entry)
XX
Cross-linking oligomer 215 fc
XX
DE Cross-linking oligomer 215 fc
XX
deoxyribonucleic acid; major
XX
deoxyribonucleic acid; major
XX
Synthetic.
XX
Synthetic.
XX
FT | Location/Qual
FT | modified_base | /*tag= a
FT | /mod_base= O'
FT | modified_base | /*tag= a
FT | modified_base | /*tag= c
FT | modified_base | /*tag= d
FT | modified_base | /*tag= d
FT | modified_base | /*tag= e
FT | modified_base | /*tag= f
FT | modified_base | /*tag= g
FT | /mod_base= m
FT | modified_base | /*tag= g
FT | /mod_base= m
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Best Local Simi
Matches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The sequence is designed to target the Human tumour necrosis beginning at nucleotide 1137 and to covalently cross-link to N4N4-ethanocytosine group. See also AAQ20031-Q20038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cross-linking oligomer 215 for targetting human TNF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New sequence-specific non-photo-activated crosslinking agents - bind to
the major groove of duplex DNA and are esp. useful for treating latent
infections e.g. HIV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         deoxyribonucleic acid; major groove; ethanoamino group;
aziridinylcytosine; cross-linking group; tumour necrosis factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-007480/01.
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14-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-MAY-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GILE-) GILEAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Match 0.3%;
Local Similarity 94.7%;
les 18; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1183 GAAAGAGAGAGAGAGAAAT 1201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAAAGAGAGAGAGAATT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 4 A; 7 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Krawczyk S;
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91US-00640654.
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                                                                                                                                                                            note= "N-methyl-8-oxo-2'-deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                     /mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "N4N4-ethanocytosine"
                                                                                                                                                                                                                                                                 'mod_base= OTHER
'note= "N-methyl-8-oxo-2'-deoxyadenine"
                                                                                                                                                                                                     base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                             base= OTHER
                                                                                                                   m5c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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RESULT 345
AAQ30384/c
ID AAQ30384 standard; DNA; 21
XX
AC AAQ30384;
XX 25-MAR-2003 (revised)
DT 25-MAR-2003 (first entry)
XX 7000 (first entry)
XX DE Oligomer TNF215 for formi)
XX Tumour necrosis factor; h
XX Tumour nec
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Best Local S
Matches 18
                                                                                                                                       Tumour necrosis factor; herpes simplex; AIDS; modified; HIV; RSV; HPV; mallgnancy; hepatitis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The sequence is designed to target the Human tumour necrosis beginning at nucleotide 1137 and to covalently cross-link to N4N4-ethanocytosine groups. See also AAQ20031-Q20038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New sequence-specific non-photo-activated crosslinking agents - bind to the major groove of duplex DNA and are esp. useful for treating latent infections e.g. {\rm HIV}.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matteucci MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 3 A; 8 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 25; 42pp; English
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14-JAN-1991;
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                                                                                                                                                                                                                 (revised)
(first entry)
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91US-00640654.
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                                                                                                                                                                                   for forming triplex with HUMTNFAA target duplex.
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/mod_base=
                                                                             Location/Qualifiers
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17
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Pred. No. 4.6e
0; Mismatches
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                    N4 N4
                  ethanocytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                               6e+02;
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The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human tumour necrosis factor beginning at nucleotide 1137 contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV, HBR, HTV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-WAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                               08-APR-1991;
17-APR-1991;
17-APR-1991;
17-APR-1991;
27-SEP-1991;
                                                                                                                                                                                                                               New oligomers contg. modified bases -doublet in a DNA duplex, for treating herpes malignancy and inflammation.
                                                                                                                                                                                                   Claim 12; Page 70; 77pp; English.
                                                                                                                                                                                                                                                                                                   WPI; 1992-217083/26.
                                                                                                                                                                                                                                                                                                                                  Froehler B,
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18-JAN-1991;
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91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
91US-00766733.
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/mod_ba
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/note= "OTHER= N4
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e= "OTHER= N6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ethanocytosine'
                                                                                                                                                                                                                                                  which form a triplex with G-C and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                    Milligan
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Sequence 21 BP; 3 A; 8 C; 0 G; 10 T; 0 U; 0 Other;

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RESULT 346
AAQ30383;
XX
AC
AAQ30383;
XX
AC
AAQ30383;
XX
DT
25-MAR-2003 (revise
DT
07-DEC-1992 (first
XX
OS
Cligomer TNF214 for
XX
OS
Synthetic.
XX
Synthetic.
XX
Synthetic.
XX
FT modified_base 1
FT modified_base 4
FT modified_base 4
FT modified_base 7
FT modified_base 7
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tumour necrosis factor; herpes simplex; AIDS; modified; HIV; RSV; HPV; malignancy; hepatitis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligomer TNF214 for forming triplex with HUMTNFAA target duplex.
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(first entry)
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                                               91WO-US008811
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/mod_base=
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/mod_base= OTHER
/mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2'
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/mod_ba
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                                                                                                                                                                         "OTHER= N4 N4 ethanocytosine"
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Pred. No. 4.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
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RESULT 347
AAT86583/c
ID AAT865
XX AAT865
XX Phosph
XX W0972:
PT wisc_c
PT
YT
YX W0972:
XX W0972
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Best Local S
Matches 18
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08-APR-1991;
17-APR-1991;
17-APR-1991;
17-APR-1991;
27-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human tumour necrosis factor beginning at nucleotide 1137 contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV, HEX, HTV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New oligomers contg. modified bases -doublet in a DNA duplex, for treating herpes malignancy and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 4 A; 7 C; 0 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; Page 70; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                         misc_difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Phosphorothioate oligonucleotide; dimeric phosphoramidite synthon; thioester; DNA synthesis; antisense oligonucleotide; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-1998
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                                                                                  06-FEB-1996;
                                                                                                                                   06-FEB-1997;
                                                                                                                                                                                          14-AUG-1997
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                              (CRUA-) CRUACHEM LTD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
91US-00766733.
                                                                                  96GB-00002326.
                                                                                                                                      97WO-GB000327.
                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
1. .21
                                                                                                                                                                                                                                                                                            /*tag= a
/note= "Phosphorothioate linkages between
nucleotides (1 and 2, 3 and 4 etc.)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  В₽
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Pred. No. 4.6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             which form a triplex with G-C and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4.6e+02;
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RESULT 348
ANA46230
ID ANA462
XX ANA462
XX ANA462
XX ANA462
XX Interp
XX Interp
KW Chromc
KW retina
KW rod- c
XX Unider
XX Claim
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Interphotoreceptor matrix; IPM; proteoglycan; IPM150; IPMC; IPM200; chromosome 6q13-q15; ocular disease; retinal detachment; chorioretinal degeneration; retinal degeneration; cone degeneration; age related macular degeneration; photoreceptor degeneration; retinal pigment epithelium degeneration; mucopolysaccharidosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence represents a phosphorothioate oligonucleotide which was prepared by solid phase synthesis. The method comprises adding at least one dimeric phosphoramidite synthon, optionally having a protected thioseter group in its internucleotide link, during the synthesis cycle. These novel dimeric phosphoramidite synthons are used as antisense molecules for inhibition of gene expression. The method gives increased yields of the phosphorothiate oligonucleotide (since fewer cycles are needed) and facilitates separation of impurities (greater difference in size compared with use of monomeric synthons)
  PCR primers AAA46209-42 were used to amplify cDNA encoding an interphotoreceptor matrix (IPM) proteoglycan, designated IPM150. Two subfamilies of IPMCs, IP protein is an IPM component (IPMC). Two subfamilies of IPMCs, IP
                                                                                                                Nucleic acids encoding interphotoreceptor matrix proteoglycans useful preventing, diagnosing and treating ocular disorders such as retinal detachment and chorioretinal degeneration.
                                                                                                                                                                                                                                                                                                                                                              29-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unidentified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA46230 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                (IOWA)
                                                                                                                                                                                                                                                                                                                       29-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200026367-A2
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                                                                                                                                                                                                                                                                                UNIV IOWA RES FOUND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    dystrophy;
                                                                                Page 44; 183pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for interphotoreceptor matrix proteoglycan IPM150 cDNA.
                                                                                                                                                                                                                                         Kuehn
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cone-rod dystrophy;
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                                                                                English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1; Length
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      IPM150
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RESULT 349
ADJ13110/c
ID ADJ1311
XX
AC ADJ131
XX
DT 20-MA;
XX
PT 20-MA;
XX
PT 14-A;
YX
YX
YX
YX
PP 27-C
XX
PP 27-C
XX
PP 27-C
XX
PP A (GA)
PP A (G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
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                                                                                                       This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in
different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequence.
                                                                                                                                                                                                                                                                                                                                                       Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                                                                             Example 1; SEQ ID NO 237; 210pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Garner HR, Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (/BEU/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    tumour suppressor; pl6; human; H69; H1618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probe; ss; chemical modification; methylation; array; CpG island;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2003152950-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1349 AAAAATTCACACAGCTGCT 1367
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MINNA J D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BALOG R P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LUEBKE K J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probe used to immobilise
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Luebke
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CpG methylated DNA SeqID 237.
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                      comprising an array of the altered DNA sequences
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RESULT 350
ADJ13108/c
ID ADJ131
XX ADJ131
XX ADJ131
XX Probe;
XW Probe;
XW Probe;
XW Lumour
XX Homo s
XX This i
CC modifi
CC invent
XX Squen
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Matches
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                                                        This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene p16 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                  Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                                                                                                                                             Example 1; SEQ ID NO 235; 210pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probe; ss; chemical modification; methylation; array; CpG island;
tumour suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human DNA probe used to immobilise CpG methylated DNA SeqID 235.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GARNER H R.
MINNA J D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LUEBKE K J.
BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GGATGAGAGGGGGAGAGGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Minna JD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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Pred. No. 4
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Sequence 21

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RESULT 351
ADJ13074/c
ID ADJ130
XX ADJ130
XX ADJ130
XX Probe;
KW Fumen
XX Probe;
KW Fumen
XX Homo 8
XX Homo 1
XX Hom
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                                                                  Query Match
Best Local S
Matches 18
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Best Local S
Matches 18
                                                                                                                                                                                                                                                 This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene p16 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                         Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (BALO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LUBB/)
                                                                                        Local Similarity
                2436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2436 GGATGAGAAGGGGAGAGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) GARNER H R.
) MINNA J D.
) LUEBKE K J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; chemical modification; methylation; array; CpG island;
suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
GGATGAGAAGGGGAGAGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGATGAGAGGGGGGAGAGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Minna JD,
                                                                                                                                                                           BP; 3 A;
                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                entry)
                                                                                           0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
                                                                                                                                                                           11 C; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Luebke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
             2454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            w
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Pred. No. 4.6e+02;
                                                                                           Score 17.4; DB 1;
Pred. No. 4.6e+02;
                                                                                                                                                                           G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ጀ
                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Balog
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.
                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21;
                                                                                                                            21;
                                                                        0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                        Gaps
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RESULT 352
ADJ1311
XX
ADJ1311
XX
DT 20-MAY
XX Probe;
KW probe;
KW tumouz
XX
Homo 8
XX
OS Homo 8
XX
OS HOMO 8
XX
PN US2003
XX
PN US2003
XX
PN (GARN)
PA (MINN)
PA (MINN)
PA (MINN)
PA (BALO)
XX
PH Garne;
XX
PH Garne;
XX
PH Garne;
XX
CT This
CC modif
CC modif
CC treat
CC inven
SQ Seque
RESULT 353
ADJ13146/c
ID ADJ131
XX
AC ADJ131
XX
DT 20-MAY
XX
DT 20-MAY
XX
DE Human
                                                                                                                                                                                                                         밁
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                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GARN/) GARNER H R.
(MINN/) MINNA J D.
(LUEB/) LUEBKE K J.
(BALO/) BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                                                                   modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analysis of chemical modification of DNA involves obtaining sample of to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
  Human DNA probe used to immobilise CpG methylated DNA SeqID 273
                                                                                                                                                                                                                                                                                                                                                                                Sequence 21 BP; 3 A; 13 C; 0 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Garner HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          14-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2003152950-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human DNA probe used to immobilise CpG methylated DNA SeqID 236.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADJ13109,
                                         20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; SEQ ID NO 236; 210pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2003-874843/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JUN-2002; 2002US-00184085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tumour suppressor; pl6; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               probe; 88; chemical modification; methylation; array; CpG island;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADJ13109 standard; DNA; 21
                                                                                ADJ13146;
                                                                                                                     ADJ13146 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-JUN-2001; 2001US-0301370P
                                                                                                                                                                                                                                                            2436 GGATGAGAAGGGGAGAGGT 2454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention relates to a novel method
                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                   18;
                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Minna JD,
                                                                                                                                                                                                                                                                                                   Conservative
                                         (first entry)
                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                      0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Luebke KJ,
                                                                                                                       21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                      Score 17.4; DB 1; Length 21; Pred. No. 4.6e+02;
                                                                                                                                                                                                                         N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            H69; H1618
                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Balog RP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for analysing chemically
                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                   0
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AAQ33
                                                                                                                                                                                                                                                                                                                                                            RESULT 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene plf from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probe; ss; chemical modification; methylation; array; CpG island;
tumour.suppressor; p16; human; H69; H1618.
                   WO9213102-A1
                                                             Bos taurus.
                                                                                                   genetic mapping;
                                                                                                                                                                 Microsatellite sequence from clone AGLA248.
                                                                                                                                                                                                           25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                                                                                                                AAQ33557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 4 A; 11 C; 0 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; SEQ ID NO 273; 210pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Garner HR, Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (LUEB/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27-JUN-2001; 2001US-0301370P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27-JUN-2002; 2002US-00184085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US2003152950-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BALO/) BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (/NINN/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (GARN/) GARNER H R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2436 GGATGAGAAGGGGAGAGGT 2454
                                                                                                   selection; primers; OPTIPRIM;
tic mapping; traits; amplificat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2003-874843/81.
                                                                                                                                                                                                                                                                                                                                                                                                                             19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MINNA J D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LUBBKB K J.
                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                             GGATGAGATGGGGAGAGGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                           (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Luebke KJ,
                                                                                                                                                                                                                                                                                                                     22
                                                                                                     amplification;
                                                                                                                                                                                                                                                                                                                     ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 17.4;
Pred. No. 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Balog RP;
                                                                                                                           breeding;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4.6e+02;
                                                                                                     88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                           cattle; parentage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
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밁 S

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RESULT 355
AAS97748
ID AAS977
XX AAS977
XX I2-MAR
DT 12-MAR
XX Murine
XX Human;
KW obesit
KW obesit
KW protei
XX D1000;
KW protei
XX W02001
XX W02001
XX W02001
XX PP 08-NOV
XX 28-APF
PR 28-APF
PR 28-APF
PR 28-JUI
PR 10-NOV
XX PA (WARN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The sequence is that of a bovine microsatellite sequence obtd. by CC screening a library of bovine MooI DNA fragments of between 250 and 500 CC bp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 CC clones cross-hybridised. Assuming independent distribution of microsatellites and MooI sites, the frequency of (T6)n >9 microsatellites CC in the bovine genome is estimated at >100, 000. The sequence information CC for ca. 230 such bovine microsatellites is summarised in the generate the creation and indexed herein (see below). The sequences upstream and CC downstream of the microsatellite sequence were used to generate the CC required PCR primers for in vitro amplification of the corresp.

CC microsatellite (using the program OPTIPRIM). The microsatellites may be considered place trait loci, or genes involved the determinism of CC consping of economic trait loci, or genes involved the determinism of CC breeding. See also AAQ33501-34437. (Updated on 25-MAR-2003 to correct PN CC field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                 Human; mouse; SAC1; carbohydrate; sweetener; ethanol; alcoholism; ss; obesity; diabetes; transgenic embryo; body tissue; body fluid; pancreas; blood; tongue; PCR primer; anorectic; antidiabetic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polymorphic bovine DNA markers - used in genetic identification, gene mapping, and selective breeding.
                                              28-APR-2000;
28-JUL-2000;
                                                                                             25-APR-2001; 2001WO-US013387.
                                                                                                                            08-NOV-2001.
                                                                                                                                                          WO200183749-A2
                                                                                                                                                                                                                      protein
                                                                                                                                                                                                                                                                                                Murine SAC1 gene-specific oligonucleotide PCR primer
                                                                                                                                                                                                                                                                                                                                                                                          AAS97748 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Table 7; Page 151; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Georges M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JAN-1991;
   (WARN ) WARNER LAMBERT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENM-) GENMARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1180
                                                                                                                                                                                                                    replacement
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AGAGAAAGAGAGAGAGA 1198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     маввеу JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.3%;
ilarity 94.7%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 11
                              2000US-0200794P.
2000US-0221419P.
2000US-0247443P.
                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91US-00642342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92WO-US000340
                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 0 C; 10
                                                                                                                                                                                                                    therapy.
                                                                                                                                                                                                                                                                                                                               entry)
                                                                                                                                                                                                                                                                                                                                                                                            22
                                                                                                                                                                                                                                                                                                                                                                                          BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0,
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             G; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  red. No. 4.76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                  #315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22;
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TCAGGCTAACCTCAAACTC 19

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Novel isolated polypeptide comprising variant form of mouse or human SAC: polypeptide, and is associated with altered preference for carbohydrates or other sweeteners, useful for preventing obesity, diabetes, alcoholism.
                                                                                                                                             Claim 14; Page 86; 239pp; English.
                                                                                                                                                                                                                            (MONE-) MONELL CHEM SENSES CENT
                                                                                                                                                                                                     Beauchamp GK,
1 DR, Ross D,
                                                                                                                                                                                                       Chatterjee
Tordoff MG;
                                                                                                                                                                                                             Þ
                                                                                                                                                                                                              De
                                                                                                                                                                                                              Jong
                                                                                                                                                                                                              PJ,
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                                                                                                                                                                                                              F.
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The invention relates to an isolated polypeptide, comprising a variant CC form of mouse or human SAC1 polypeptide. The variant form is associated CC with altered preference for carbohydrates, other sweeteners or ethanol. CC The polypeptide and its associated DNA sequence can be produced by CC recombinant techniques and is useful for preventing obesity, diabetes or CC alcoholism associated with SAC1 expression. The sequences are useful in CC embryos may be used in screening for and identifying agents that induce CC embryos may be used in screening for and identifying agents that induce CC embryos may be used in screening for and identifying agents that induce CC embryos may be ascertained by testing any fluid or tissue of a human CC (such as blood, pancreas or tongue) for sequence variations of the SAC1 (cgene. A sequence variation of the SAC1 locus may indicate a CC diagnostic mark. The polymucleotide can be detected in a biological CC sample by contacting the DNA with a probe to form a hybridisation complex CC which is then detected. The sequences represent cDNA encoding human and CC mouse SAC1 polypeptides and PCR primers specific for the SCA1 genes Sequence 22 BP; 8 A; 8 C; 2 G; 4 T; 0 U; 0 Other;

Query Match Best Local S Matches 18 3597 TCAGGCTAATCTCAAACTC 3615 18; Similarity Conservative 0.3%; 94.7%; ° Score 17.4; Pred. No. 4. Mismatches 4.7e+02 DB 1; Length 22; <u>,,</u> Gaps 0;

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09-JUL-2003
                   ACA89735
                                      ACA89735 standard;
(first entry)
                                       DNA;
                                       22
                                       ₽P
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Herbicide resistance polymorphic marker related primer #34.

glyphosate; paraquat; sulfonyl urea Polymorphic marker; herbicide resistance; herbicide susceptible plant; herbicide resistant plant; Conyza canadensis; Lolium rigidum; goosegrass; moiety; PCR; primer;

RESULT 356
ACA89735
ID ACA8977
XX ACA897
XX O9-JUL
XX POlymc
KW Polymc
KW Polymc
KW 191yphc
XX Synthe
XX Synthe
XX I-OCI
XX I-OCI WPI; 2003-430273/40. 12-OCT-2001; 2001US-0328750P 11-OCT-2002; 2002WO-US032637. 17-APR-2003 WO2003031937-A2 Ó MORPHOTEK INC Grasso L, Nicolaides Š, Sass

Identifying polymorphic

markers

of herbicide resistance

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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention describes a method of identifying polymorphic markers of herbicide resistance in a plant. The method involves: isolating genomic DNA from an herbicide susceptible plant and an herbicide resistant plant of the same species, performing genetic analysis and identifying the difference that correlate with herbicide resistance DNA, identifying the difference that correlate with herbicide resistance or susceptibility, thus identifying polymorphic markers. The method is useful for identifying polymorphic markers of herbicide resistance in a plant e.g. Conyza canadensis, Lolium rigidum and goosegrass species, where the herbicides include glyphosate, paraquat and sulfonyl urea moieties. This sequence represents a primer associated with the identification of polymorphic markers of herbicide
                                                                                       Bachmanov
Ross DA,
          or ethanol, us
gene function
            New isolated polynucleotides for sensing carbohydrates, other sweeteners or ethanol, useful for screening drugs for inhibition or restoration of gene function as antidiabetic, antiobesity or antialcohol consumption
                                                                                                                                                                              (REED/
                                                                                                                                                                                                                                                                                                                                                   US2004081964-A1
                                                                                                                                                                                                                                                                                                                                                                                                                Mouse; SAC1; PCR; ss; carbohydrate; sweetener; ethanol; obesity; diabetes; alcoholism; antidiabetic; alcohol; anorectic; antialcoholic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Murine SAC1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADM16088;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADM16088 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22
                                                                                                                                                                                                                                                                       25-OCT-2002; 2002US-00280183
                                                                                                                                                                                                                                                                                                25-OCT-2002; 2002US-00280183.
                                                                                                                                                                                                                                                                                                                                                                             Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          analyzing genomic DNA of herbicide resistant and susceptible plants, and identifying difference that correlate with resistance or susceptibility.
                                                                                                                                                       /WHO
                                                                                                                                                                                                                     LISS/
                                                                                                                                                                                                                                  (BEAU/)
                                                                                                                                                                                                                                             BACH/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1180 AGAGAAAGAGAGAGAGAA 1198
                                                               2004-340133/31.
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                                                                                                                           ROSS D A.
OHMAN J D.
CHATTERJEE A.
DE JONG P J.
                                                                                                                                                                             REED D R.
TORDOFF M G.
                                                                                                                                                                                                        LI S.
                                                                                                                                                                                                                                 BACHMANOV A
BEAUCHAMP G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGAGAGAGAGAGAGAGA 20
                                                                                     AA, B
Ohman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                    Beauchamp GK, Li
n JD, Chatterjee
                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primer #315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10 A; 0 C; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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Pred. No. 4.7e+02
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0
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                                                                                       Li X,
De Jong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 22;
                                                                                        Reed
PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Indels
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The invention relates to SAC1 polypeptides and the polymucleotides CC encoding them. The polymucleotides contain a variation associated with CC sensing carbohydrates, other swetteners or ethanol. The invention also CC relates to a method for analysing a biomolecule in a biological sample, CC comprising altering SAC1 activity in the sample and measuring the CC comprising contacting a polymucleotide in a biological sample, CC comprising contacting a polymucleotide in a biological sample with a CC probe where the probe hybridises to a SAC1 polymucleotide to form a CC hybridisation complex and detecting the hybridisation complex, a method CC of identifying susceptibility to obsaity or diabetes comprising comparing the nucleotide sequence of the suspected SAC1 allele with a wild type CC nucleotide sequence, where the difference between the suspected allele CC and the wild-type sequence identifies a sequence variation of the SAC1 comprising CC diabetes or alcoholism associated with expression of SAC1, comprising CC administering to a subject a pharmaceutical composition and a transgenic CC animal that carries an altered SAC1 allele. The methods and compositions of the invention are useful for screening drugs for inhibition or CC consumption therapies and for identifying sweeteners and alcohols. This sequence represents a PCR primer used to amplify murine SAC1 DNA of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 12;
                                                      invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 358; 148pp; English.
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Query Match Best Local S Matches 18 Local Similarity 3597 TCAGGCTAATCTCAAACTC 3615 18; -TCAGGCTAACCTCAAACTC 19 Conservative 0.3%; 0, Score 17.4; DB 1; Pred. No. 4.7e+02; Mismatches Length Indels 22 0 Gaps

0

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Sequence 22

BP; 8 A; 8 C; 2 G; 4 T; 0 U; 0 Other;

RESULT 358
AAT13515
ID AAT135
XX
AC AAT135
XX
DT 20-NOV
XX
DE Telome
XX
Human
KW Human
KW interi
KW telome
XX
Homo
DPN GB229
XX
DF 19-OC
XX
PF 19-OC
PR 21-OC
PR 21-O GB2294322-A Telomere mapping primer (TelG) for Xp/Yp PAR1 proximal target AAT13515 standard; telomère mapping; human chromosome; cancer identification; ss. internal repeat unit; detection; disease predisposition; abnormal; cell growth; cell division; diagnosis; family relationship; allele primer; 20-NOV-1996 AAT13515; sapiens. Xp/Yp; (first entry) pseudoautosomal region; PAR1; telomere repeat array;
; genomic DNA; characterisation; amplification; genomic DNA; characterisation; DNA; 23 ВP

19-OCT-1995; 24-APR-1996 95GB-00021447

21-OCT-1994; 21-MAY-1995; 94GB-00021234. 95GB-00010639.

(ZENB) ZENBCA LTD.

묫 Royle 'n, Jeffreys 2

WPI; 1996-190531/20.

Characterising genomic DNA by in telomere repeat array used amplifying selected internal repeat units - for, e.g. disease diagnosis, determn. of

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RESULT 359
AAL57030/c
ID AAL570
XX AAL570
XX AAL570
AC AAAL570
AC AAL570
AC AAAL570
AC AAACA
AC 
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A sequence proximal to the human Xp/Yp pseudoautosomal region (PAR1) telomere repeat array, was produced using sequence information obtd. from 32 Caucasian and 21 African DNAs. The sequence was found to contain 20 substitution polymorphisms, and a 10 bp deletion/insertion polymorphism found only in African DNAs. Claimed type specific primers, which hybridise to loci contained in the sequence, can be used for the characterisation of a genomic DNA test sample by amplifying selected internal repeat units in a telomere repeat array. This characterisation is useful in the detection of inherited or acquired disease predisposition, abnormal cell division or growth diagnosis (e.g. cancer), individual and family relationship identification and telomere mapping of thuman chromosomes from DNA flanking sequences, i.e. 2 state mapping using
The present invention relates to a polypoptide comprising vascular endothelial-protein tyrosine phosphatase (VE-PTP) or its active fragment or effector, or the nucleic acid encoding the polypoptide or its effector, for use in the manufacture of an agent for monitoring or modulating VE-cacherin mediated processes or disorders. The polypoptide comprising vascular endothelial-protein tyrosine phosphatase (VE-PTP) or its active fragment or effector, or the nucleic acid encoding the
                                                                                                                                                                                                                                                                                                                                                                                        Use of the polypeptide comprising vascular endothelial-protein tyrosine phosphatase (VE-PTP) or the nucleic acid encoding the polypeptide for the phosphatase of an agent for monitoring or modulating VE-cadherin mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vascular endothelial protein-tyrosine phosphatase; VE-PTP; gene therapy; cytostatic; VE-cadherin; PCR; primer; ss; vascular endothelial-cadherin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nawroth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           08-APR-2002; 2002EP-00007837
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ₽,
                                                                                                                                                                                                                                                                                     Page 17; Opp; English.
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Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Golding
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                                                                                                                                                                                 active fragment
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                                        ဝူ
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문 S

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                                               Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 19
                                                                                                                                 The present invention relates to a panel of two or more single nucleotide polymorphisms, where each of the polymorphisms of the panel are selected from single nucleotide polymorphisms that are not genetically linked with respect to one another, and where each of the polymorphisms of the panel are selected from single nucleotide polymorphisms that are located outside tandem repeat nucleic acid sequences. The known sample is from a family member. The compromised nucleic acid sequences comprises nucleic acid fragments from 10-100 nucleotides in length. The identity of the one or more single nucleotide polymorphisms is determined using a single base primer extension reaction. The present sequence represents a primer of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polypeptide or its effector, is useful for the manufacture of an for monitoring or modulating VE-cadherin mediated processes or die.g., cancer. The present sequence is a PCR primer shown in the
                                                                                                Sequence 24
                                                                                                                                                                                                                                                                                                                              New panel of single nucleotide polymorphisms comprising two or more single nucleotide polymorphisms, useful for analyzing compromised nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADO17957;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADO17957
                                                                                                                                                                                                                                                                                         Disclosure; SEQ ID NO 184; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-091088/09
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Primer of the invention #183.
                                                                                                                                                                                                                                                                                                                                                                                                                    (ORCH-) ORCHID BIOSCIENCES INC
                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JUN-2002; 2002US-0392504P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-JUN-2003; 2003WO-US020150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO2004003220-A2
                                                             Local Similarity
                        1182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5392 TAAAAAAATACAAAAAAGAAAAA 5414
                                                                                                                                                                                                                                                                                                                  sampies.
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                                                 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleotide polymorphism; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                        AGAAAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
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AGAAAGAGAGAGAAAAA
                                                                                                BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 0 A; 0 C; 0
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                                                 Conservative
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                                                                                                                                                                                                                                                                                                                            polymorphisms,
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                                                          0.3%;
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                                                                                                  10 C; 0
                                                                                                                                                                                                                                                                                                                                                                                           Mckeown B,
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                                                0;
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Pred. No. 4.7e
1; Mismatches
                                                                                                G; 14
                                                             Score 17.4; DB Pred. No. 4.7e+(
                                                 Mismatches
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                                                              4.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 Other;
                                                                                                   0 Other;
                                                                         Length 24;
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                                                 Indels
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                                                 Gaps
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RESULT 361
AAQ64724
                                                       Query Match
Best Local Similarity
Matches 19; Conserv
                                                                              This sequence was used to determine whether 2-5A-antisense chimeric molecules are inhibitory to cell growth. The molecules AAQ64701, AAQ64771 and AAQ64724 all lacked cytotoxicity. In the movel 2-5A-antisense oligonucleotide chimeric molecules, the antisense region targets the chimeric molecule to a particular region of RNA to be specifically cleaved and the 2',5'-linked tetraadenylate tail activates the 2-5A RNase. Typical applications are treatment of viral infections (esp. for cleavage of an RNA virus genome), cancer; leukaemia, cardiovascular disorders (e.g. restenosis after angioplasty), genetic disorders, osteoarthritis or rheumatoid arthritis. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                              Specific cleavage of RNA, useful partic. for treating viral infection, cancers, etc. - by using anti-sense oligo:nucleotide coupled to activator of 2-5A dependent RNase.
                                                                                                                                                                                                                                                                                                                                                                                           (USSH )
                                                                                                                                                                                                                                                                                                                                                                                                                                     21-OCT-1992;
17-SEP-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein synthesis regulation; phosphorylation; eIF-2alpha;
eukaryotic translation initiation factor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2003
04-JAN-1995
                                                       Sequence 22
                                                                                                                                                                                                                                                     Example 9;
                                                                                                                                                                                                                                                                                                                                     WPI; 1994-151315/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-OCT-1993,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-APR-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2',5'-linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ64724 standard; cDNA to mRNA;
                                                                                                                                                                                                                                                                                                                                                               Torrence P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9409129-A2
                                                                                                                                                                                                                                                                                                                                                                                           US DEPT HEALTH & HUMAN SERVICES. CLEVELAND CLINIC RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2',5'-tetraadenylate; 2-5A dependent RNase activator;
ge; antiviral therapy; chimeric molecule; PKR;
                                                                                                                                                                                                                                                    Page 66; 86pp;
                                                     B₽;
 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tetraadenylate-anti(dT)18 oligonucleotide chimeric mol.
                                                                                                                                                                                                                                                                                                                                                               Silverman R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antiviral therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                    92US-00965666.
93US-00123449.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93WO-US010103
                                                     22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "the 2-5A moiety (*tag = a) and the antisense
sequence (*tag = c) are linked by two 1,4-butanediol
molecules linked through phosphodiester bonds"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= a
/label= 2',5'-linked tetraadenylate
/note= "nucleotides linked through phosp)
/note= "aroups of 2' and 5' carbons"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                     ₽,
             0.3%;
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                                                     C; 0
                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                               Maitra R,
 <u>.</u>
Score 17.2; DB 1; Length Pred. No. 5e+02; Indels 0; Mismatches 3; Indels
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                                                     0 Other;
 Indels
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 Gaps
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RESULT 362
AAF17413
ID AAF174
XX AAF174
XX OP-MAR
XX PT 09-MAR
XX Retrot
XX Uniden
XX 
RESULT 363
ADF12348
ID ADF123
XX ADF123
XX ID 12-FEB
DY 12-FEB
DX L1 ret
XX Gene t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to DNA for a promoter and an L1 cassette sequence having a core retrotransposon element. The invention is useful for random insertion of a heterologous or homologous DNA sequence into a cell genome, and for correction of a genetic defect in the cell into which the insertion is made. Genetic defects which may be corrected includes cystic fibrosis, mutations in the dystrophin gene, genetic defects associated with blood clotting and other genetic defects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5393 AAAAAAATACAAAAAAGAAAAA 5414
  gene therapy; insertional mutation; germ line specific promoter;
                                                  L1 retrotransposon endonuclease cleavage site seq id
                                                                                                       12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 22 BP; 22 A; 0 C; 0 G; 0 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNAc comprising a promoter P and an L1 cassette sequence having a core retrotransposon element, useful for random insertion of a heterologous homologous DNA sequence into a cell genome and for correcting genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-060015/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-NOV-1995;
15-NOV-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Retrotransposon;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    L1 cleavage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-MAR-2001
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                                                                                                                                                                                                             ADF12348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-APR-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          l Similarity
19; Conserv
                                                                                                                                                                                                             standard;
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                                                                                                                                                                                                                                                                                                                                       AAAAAAAAAAAAAAAAA 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dombroski BA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fig 14; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       site related sequence #3
                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-0006831P
96US-00749805
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genetic defect; cystic fibrosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.3%;
                                                                                                       entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 17.2;
Pred. No. 56
                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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RESULT 364
ADQ25630/c
ID ADQ256
XX A
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention describes a method of creating an insertional mutation in the germ line of an animal by introducing into an animal a nucleic acid molecule comprising a germ line specific promoter. The method is useful for generating a mutation in an offspring of an animal, or for isolating a nucleic acid from a genome of an offspring of an animal. The method may also be used to correct genetic defects in animals, especially humans. The nucleic acid is useful for generating mutations in a cell for assessing the frequency with which selected cells under go insertional mutagenesis for the generation of transgenic animals. This sequence represents an exemplary cleavage site of the endonuclease encoded by human L1 retrotransposon EN domain.
                                                                                                                                                                         Cystic fibrosis; muscular dystrophy; diabetes; gene discovery; gene mapping; molecular haplotyping; agricultural research; prostate cancer; breast cancer; lung cancer; colon cancer; ovarian cancer; human; adenorectal carcinoma; primer; ss.
                                                                US2004126770-A1
                                                                                                                                                                                                                                                                                                                             Junction-specific poly(A) cDNA primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADQ25630 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 22 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Creating an insertional mutation in the germ line of an animal, useful for generating a mutation in an offspring of an animal, comprises introducing into an animal a nucleic acid molecule comprising a germ line
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01-SEP-2000;
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                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                                                                              23-SEP-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        16-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-AUG-2002; 2002US-00216122.
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96US-00749805.
97US-00847844.
2000US-00653812.
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KW Anorec
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    identifying or analysing and comparing RNA molecules and or sequences expressed in different cells, tissues and or samples. The invention is also useful in detecting disease (e.g. cystic fibrosis, muscular dystrophy or diabetes), mutation detection, gene discovery, gene mapping (molecular haplotyping), agricultural research, and assessment of predisposition for cancers, e.g. prostate, breast, lung, colon or ovariar cancer. The present sequence is a junction-specific CDNA primer. This sequence is used to illustrate the method of invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to composition and method for amplifying RNA sequences. The method involves synthesising first strand cDNA molecules from RNA molecules, circularising the first strand and replicating the circularised first strand cDNA molecules using rolling circle replication. The method is useful for producing nucleic acid molecules corresponding to RNA molecules in an RNA sample, for
                                                                                                                                                                                                                                                                   Anorectic; Antidiabetic; Muscular; Gene Therapy; CpG island; IGF2 gene intron 3; muscle mass; fat deposition; teat number; obesity; muscle deficiency; diabetes; SNP; single nucleotide polymorphism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 0 A; 0 C; 0 G; 22 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADQ80747 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Amplifying RNA sequences, useful in detecting diseases or mutation, comprises synthesizing first strand cDNA, circularizing first strand cDNA molecules by rolling cicDNA, and replicating the circularized cDNA molecules by rolling ci
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                                                                      14-JUL-2004
                                                                                                          EP1437418-A1
                                                                                                                                                                                 variation
                                                                                                                                                                                                                                     Sus scrofa.
                                                                                                                                                                                                                                                                                                                                            Porcine KVLQT1 intron 12 DNA sequence polymorphism oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                 23-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; SEQ ID NO 6; 64pp; English.
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(ABAR/) ABARZUA P.
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ilarity 86.4%;
Conservative
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                                                                                                                                                                         Location/Qualifiers replace(13,T)
                                                                                                                                                              /*tag=
                                                                                                                                          standard_name= "Single_nucleotide_polymorphism"
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Pred. No. 56
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10-JAN-2003; 2003EP-00075091.

10-JAN-2003; 2003EP-00075091.

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RESULT 366
AAQ30430/c
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IGF2 gene by modulating the activity of an evolutionarily conserved CpG island located in intron 3 of an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene. Also claimed is a method (M3) for identifying a compound capable of modulating mRNA transcription of an IGF2 gene and a method (M4) for identifying a compound capable of modulating binding of a nuclear factor to an IGF2 gene. (M2) is useful for modulating mRNA transcription of an IGF2 gene in a cell or organism. (M3) and (M4) are useful for identifying compounds capable of modulating mRNA transcription of an IGF2 gene and/or modulating binding of a nuclear factor to an IGF2 gene. Compounds identified are potentially useful for treating obesity, muscle deficiencies and diabetes. The present sequence is a porcine sequence tagged sites (STS) comprising a DNA sequence polymorphism, which was isolated in an example from the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a method (M1) for selecting an animal for having desired genotypic or potential phenotypic properties. (M1) comprises testing the animal for the presence of a nucleic acid modification affecting: the activity of an evolutionary conserved CpG island located in intron 3 of an IGF2 gene; and/or binding of a nuclear factor to an IGF2 gene. The nuclear factor is capable of binding to a stretch of nucleotides which in the wild type pig, mouse or human IGF2 gene is part of an evolutionary conserved CpG island, located in intron of the IGF2 gene. The stretch is functionally equivalent to (ADQ80709). The nucleic acid modification in ADQ80709 comprises a G to A transition at IGF2-intron3-nt3072. (M1) is useful for selecting an animal with properties related to muscle mass, fat deposition, and/or teat number. Also claimed is a method (M2) for modulating mRNA transcription of an animal with the selecting manufactor of an animal with the selecting manufactor of an animal with the selecting mRNA transcription of an animal with the selecting m
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                                                                                                                                                                                                 Human interleukin-6 gene; herpes simplex; AIDS; modified; HIV; RSV; malignancy; hepatitis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
07-DEC-1992
                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                    Oligomer IL6803 for forming triplex with HUMIL6 target duplex.
                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ30430;
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                                                              modified_base
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                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
                                                                                        Location/Qualifiers
/*tag= a
/mod_base= OTHER
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Pred. No. 5
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RESULT 367
AAQ30431/c
ID AAQ304:
XX
AC AAQ304

standard; DNA; 23

AAQ30431; AAQ30431 맑 S

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The synthetic oligomer is capable of forming a triplex at physiological CC pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human interleukin 6 gene untranslated sequence contg. a purine rich sequence CC concd. on one strand of the duplex. The oligomer, and others like it are CC useful in diagnosis and therapy of diseases characterised by specific DNA CC duplex targets, e.g. HPV, HEW, HIV, hepatitis B, herpes, malignant CC thus assays may be carried out without subjecting the test specimen to thus assays may be carried out without subjecting the test specimen to CC thus assays may be carried out without subjecting the test specimen to CC formed from an o-xyloso dimer synthon. The linking gp. is o-xyloso CC (nucleotides have the 3'positions of xylose sugars linked via the o-CC xylene ring). Two nucleotides are coupled through a xylene residue to CC cycle ring. The oligomer is able to inhibit colligomer stable to nuclease activity. The oligomer is able to inhibit colleges and an accoupled a cycles and Angolaca And (malated on Schwarzona to correct by field)
                                                                                            Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New oligomers contg. modified bases - doublet in a DNA duplex, for treating herpes malignancy and inflammation.
                                                                                                                                                                                                               Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 12; Page 71; 77pp; English.
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27-SEP-1991;
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18-JAN-1991;
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17-APR-1991;
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                                               5395 AAAAATACAAAAAAGAAAAAT 5416
22
                                                                                                                                                                                                            BP; 2 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                       Conservative
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91US-00686546.
91US-00686547.
91US-00766733.
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91US-00643382.
91US-00683420.
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/note= "OTHER= N6 methyl-8-oxo 2' deoxyadenine"
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/note= "o-xyloso dimer synthon linkage"
12. 23
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/note= "see comments"
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                                                                                                    0; Mismatches
                                                                                                                                 Score 17.2; DB Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       which form a triplex with G-C and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Milligan
                                                                                                                                                          DB 1; Length 23;
                                                                                                                                                                                                                                                                       to correct PN field.)
                                                                                                          Indels
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                                                                                                          Gaps
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The synthetic oligomer is capable of forming a triplex at physiological CC pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human cC interleukin 6 gene untranslated sequence contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are cc useful in diagnosis and therapy of diseases characterised by specific DNA CC duplex targets, e.g. HPV, HER, HTV, hepatitis B, herpes, malignant cc thumours and inflammation. The triple helices form under mild conditions the conditions. The oligomer contains an inverted polarity region cc thus assays may be carried out without subjecting the test specimen to charsh conditions. The oligomer contains an inverted polarity region cc formed from an o-xyloso dimer synthon. The linking gp. is o-xyloso cc (nucleotides have the 3 positions of xylose sugars linked via the o-cc xylene ring). Two nucleotides are coupled through a xylene residue to oligomer stable to nuclease activity. The oligomer is able to inhibit cc gene expression, as verified by in vitro systems. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-APR-1991;
17-APR-1991;
17-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-NOV-1990;
18-JAN-1991;
08-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 12; Page 71; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New oligomers contg. modified bases -doublet in a DNA duplex, for treating herpes malignancy and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             25-NOV-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligomer IL6804 for forming triplex with HUMIL6 target duplex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
07-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GILE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GILEAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Β,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Krawczyk S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
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91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
91US-00766733.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "see
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= c
/label= inverted_polarity_region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= d
/note= "o-xyloso dimer synthon linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11. .12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           base= OTHER
= "OTHER= N6 methyl-8-oxo 2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 _base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      "OTHER=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       σ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matteucci MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comments"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N4 N4 ethanocytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    which form a triplex with G-C and diagnosing HIV, hepatitis,
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RESULT 369 AAF16627/c

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Best Local Similarity
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Best Local Similarity
Matches 19; Conserv
                                                       Matches
                                                                                                                                    The specification describes a process for circularizing an oligonucleotide around a double-stranded nucleic acid that contains a target sequence. The method is used to detect or label nucleic acids, particularly plasmids, to detect target sequences in the nucleic acid, and to distinguish between two sequences that differ in only 1 or 2 mutations. It can be used to select, e.g. from degenerate single-stranded nucleic acids, sequences that can bind to the nucleic acid, particularly sequences that promote entry of the nucleic acid into cells or can target the nucleic acid to specific cellular compartments. The method can also be used to purify nucleic acids, particularly plasmids, and in gene therapy for specific inhibition of a gene contained in the nucleic acid. The present sequence represents an oligonucleotide used in the course of the invention, during construction of a triple helix
                                                                                                            Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                 Circularizing oligonucleotide around double-stranded nucleic acid, usefie.g. for detecting mutations, using target-binding oligonucleotide with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid detection; nucleic acid labelling; gene therapy; nucleic acid purification; triple helix; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABL56769;
                                                                                                                                                                                                                                                                                                                                           Example 10;
                                                                                                                                                                                                                                                                                                                                                                     complementary end sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Escude C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-JUN-2000; 2000WO-FR001655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200077250-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence of an oligonucleotide used for triple helix constrution
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABL56769 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (INRM ) INSERM INST NAT SANTE & RECH MEDICALE. (CNRS ) CNRS CENT NAT RECH SCI.
                          2560 GATGAGGGGAGAGAGAGATGG 2581
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 GAAGAGGGAGAGAGAGAGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Garestier T,
                                                                                                                                                                                                                                                                                                                                         Page 40; 91pp;
                                                       Conservative
                                                                                                              BP; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99FR-00007503.
                                                                                                              A; 0 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
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                                                                                                              12 G; O T;
                                                                                                                                                                                                                                                                                                                                           French.
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                                                                   Score 17.2; DB Pred. No. 5e+02;
                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Roulon
                                                                                                              0 U; 0 Other;
                                                                                 DB 1; Length
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                                                      Gaps
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RESULT 370
ABLO1773
ID ABLO17
XX ABLO17
XC ABLO17
XX IS-MAR
XX IS-MAR
XX Human
XX Human;
KW Human;
KW Human;
KW Human;
KW Horedi
XX Homo &
XX IS-MON &
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention provides oligonuclectides, and methods for their use, which are useful in modulating the action of proteins involved in gastric acid production. The target protein is preferably the histamine H2 receptor or one of the proteins which form part of the gastric proton pump. The sequences and methods of the invention are useful in the treatment of gastric reflux, gastritis, dyspepsia, stomach ulcers, duodents ulcers and other gastric acid disturbances, most of which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gastric acid disturbance; gastric reflux; gastritis; dyspepsia; stomach ulcer; duodenal ulcer; Helicobacter pylori; antisense; DNA-RNA hybrid; ss.
                                                                                                                                                                                                Human; MLH1; MSH2; hMLH1; hMSH2; variant gene; diagnosis; HNPCC; hereditary non-polyposis colorectal cancer; ds.
                                                                                                                                                                                                                                                                                         Human MSH2
                                                                                                                                                                                                                                                                                                                                                 18-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 23 BP; 1 A; 0 C; 0 G; 22 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            production or secretion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating gastric acid disturbance by administering an oligonucleotide which modulates the activity of a polypeptide involved in gastric acid
                                22-NOV-2001
                                                                                     US2001044936-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL01773 standard; DNA; 23 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          caused by Helicobacter pylori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2001-025093/03
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gastric acid production inhibiting oligonucleotide SEQ ID NO: 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-MAR-2001
                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (TACH/) TACHAS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
Lee 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5395 AAAAATACAAAAAAGAAAAAAT 5416
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                                                                                                                                                                                                                                                                                   (hMSH2) intronic sequence SEQ ID NO:126.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                              (first entry)
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Pred. No. 5e+02;
0; Mismatches
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ARESULT 371
ARESULT 371
ARESULT 112/C
ID AAL571
XX ARESULT
AC AAL571
XX Human
XX Human
XX Human
XX Human
XX BB.
XX BB.
XX BOZ 103
XX BZ
PN WOZ 103
XX ZZ
PD 22-WAY
XX BR.
XX 15-NON
XX PP 15-NON
XX PR 16-NON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a variant human MLH1 or MSH2 gene. Also Cd described are: (1) a method for diagnosing or predicting susceptibility to hereditary non-polyposis colorectal cancer (HNPCC), comprising CC screening a DNA sample for the variant MLH1 or MSH2 gene where presence of the variant indicates presence of, or susceptibility to HNPCC; (2) a CC method of identifying mutants in splice donor or acceptor sites of a human MLH1 gene, comprising sequencing splice donor or acceptor sites of the gene with intronic primers for the human MLH1 gene and analysing the CC sequence to identify any mutants; (3) a method of identifying mutants in splice donor or acceptor sites of sequencing splice donor or acceptor sites of the gene with intronic comprising splice donor or acceptor sites of the gene with intronic comprising splice donor or acceptor sites of the gene with intronic comprising splice donor or acceptor sites of the gene with intronic comprising calls expressing the variant MLH1 or MSH2 gene. The hMLH1 and CC susceptibility to hereditary non-polyposis colorectal cancer. ABL0145 and ABL01745 to ABL0184 to ABL01831 represent invention. ABL01831 represent invention of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                        15-NOV-2002; 2002WO-IT000729
                                                                                                            22-MAY-2003
                                                                                                                                               WO2003042409-A2
                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                     Human epithelial caderine; E caderine; gastric carcinoma; PCR; primer;
                                                                                                                                                                                                                                                                         Human epithelial caderine PCR primer 2 (from primer pair A).
                                                                                                                                                                                                                                                                                                            17-SEP-2003
                                                                                                                                                                                                                                                                                                                                              AAL57112;
                                                                                                                                                                                                                                                                                                                                                                                 AAL57112 standard; DNA; 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 23 BP; 21 A; 0 C; 1 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New variants of the human MLH1 and MSH2 genes for diagnosing or determining a predisposition for hereditary non-polyposis colorectal
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                                     16-NOV-2001; 2001IT-TO001077
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(LING/) LIN-GOBRKE J L.
(LING/) LING J C.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5392 TAAAAAATACAAAAAAGAAAA 5413
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19; Conserv
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Pred. No. 5e
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(UYUR-) UNIV URBINO

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RESULT 372
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      lymphokine; heparin binding growth factor; proliferation; primer; collagen synthesis; hyaluronan synthesis; chemoattractant; amplify; wound healing; antibody fibrosis; angiogenesis; inflammation; PCR; liver cirrhosis; primary biliary cirrhosis; glomerulnephritis; renal failure; adult respiratory distress syndrome; cystic fibrosis; asthma; emphysema; pulmonary fibrosis; gastrointestinal fibrosis; Crohn's disease; ulcerative colltis; intestinal occlusion; fibrotic cancer; optic fibrosis; dermal fibrosis; scleroderma; marrow fibrosis; joint fibrosis; vascular fibrosis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel method for the diagnosis of greater susceptibility to gastric carcinoma, comprising searching for a possibl polymorphism in the promoter of the epithelial caderine (E-caderine) gene. The method is useful for identifying a genetic polymorphism that leads to a greater predisposition to the onset of gastric carcinoma. The method is relatively simple, quick, accurate and reliable. The present sequence is that of E-caderine PCR primer 2 (from primer pair A) used during a method to identify the genotype of an individual for a C to A polymorphism at nucleotide -160 of the E-caderine gene and claimed in claim 11 of the specification
                               Fibroblast stimulating pathological scarring of fibrosis etc.
                                                                                                                                                                                                                                                                                                          28-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                      06-SEP-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Murine fibrosin 2B3 domain derived primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAT33605
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                                                                                                                                                                                                                                                 (NEWE-) NEW ENGLAND MEDICAL CENT
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                                                        g factor-1 caused by
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Pred.
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                                                        polypeptide - used in treatment
inflammation and for suppression
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                                                                                                                                                                                                                                                       HOSPITALS INC.
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                                                   of
of
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RESULT 373
AAT61390/c
ID AAT613
XX A
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CC Fibrosin may be used in therapeutic compositions to stimulate wound chealing. Antibodies against fibrosin may be used for suppressing chealing. Antibodies against fibrosin may be used for suppressing chiflammation. These methods may be used in patients which have been chiflamy cirrhosis, glomerulnephritis, renal failure, adult respiratory clistness syndrome, cystic fibrosis, asthma, emphysema, pulmonary chiflers, gastrointestinal fibrosis, Crohn's disease, ulcerative colitis, intestinal occlusion, a fibrotic cancer, optic fibrosis, dermal fibrosis,
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Best Local S
The sequences given in AAT61387-91 are a probe and primers which used in the isolation of full length cDNA's encoding murine and h fibroblast stimulating factor (FsF)-1, also known as fibrosin. Fi
                                                                                                                                                                                  Fibroblast stimulating factor (FsF)-1 polypeptide - used in the detection of propensity for pathological fibrosis, esp. resulting from sarcoidosis, scleroderma, rheumatoid arthritis etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       lymphokine; heparin binding growth factor; proliferation; collagen synthesis; hyaluronan synthesis; chemoattractant; wound healing; antibody fibrosis; angiogenesis; inflammation; hepatic fibrosis; schistosomiasis; sarcoidosis; scleroderma; sclerosing cholangitis;
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                                                                                                                                                                                                                                                                                                                                                                                 Wyler DJ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Murine; human; fibroblast stimulating factor; FsF-1; fibrosin;
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                                                                                                                           Example; Page 53; 144pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          rheumatoid arthritis; ulmonary fibrosis; intestinal
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                                                                                                                                                                                                                                                                                                                                                                                 Prakash S,
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Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  HOSPITALS
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RESULT 374
AAA448/C
ID AAA4482
XX AA4482
XX 1-SEP
XX 1-SEP
XX 1-SEP
XX Heart
XX Heart
XX HOMO 8
XX POR DT
OS-JUL
XX 27-DEC
XX 28-DEC
PR 28-PEB
PR 02-NOV
XX (KYOW
XX UMEZAW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                       The present invention provides cells originating in the human bone marrow or umbilical blood cells which are capable of differentiating into cardiomyocytes. These cells are useful in the treatment of diseases involving heart muscle degeneration, such as myocardial infarction, and the study of cardiomyocyte differentiation. The present sequence is a PCR primer described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            is a lymphokine which is a heparin binding growth factor. It stimulates fibroblast proliferation, collagen and hyalinronan synthesis and acts as a chemoattractant for fibroblasts. The 283 domain exhibits comparable activity to full length fibrosin in a cellular proliferation assay. Fibrosin may be used in therapeutic compositions to stimulate wound healing. Antibodies against fibrosin may be used for identifying individuals with a propensity for pathological fibrosis, esp. hepatic fibrosis. This method may be used on patients who are thought to be suffering from schistosomiasis, or where the pathological fibrosis results from sarcoidosis, scleroderma, sclerosing cholangitis, rheumatoid arthritis, pulmonary fibrosis or intestinal pneumonitis
                                                                                                                                                                                                                                                                                                                                                                                          Cells capable of differentiating into cardiomyocytes and originating bone marrow or umbilical blood cells for study of cardiomyocyte differentiation and treatment of heart disease.
                                                                                                            Sequence 24
                                                                                                                                                                                                                                                                                                                                               Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-DEC-1999;
28-FEB-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Heart muscle cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-425656/45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yamada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           27-DEC-2000; 2000WO-JP009323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200148151-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Heart muscle cell; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 24 BP; 2 A; 11 C; 2 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Omezawa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (KYOW ) KYOWA HAKKO KOGYO KK.
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                                                                                                                                                                                                                                                                                                                                          Page 156; 183pp; Japanese.
          Conservative
                                                                                                            BP;
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2000WO-JP007741.
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                                0.3%;
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                                                                                                            7 C; 4 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Fukuda
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell differentiation; heart disease;
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                             Score 17.2; DB 1; Pred. No. 5e+02;
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          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ogawa
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                                                      Length
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        Indels
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RESULT 375
AAH49612/C
ID AAH496
XX AAH496
XX beta-s
XX beta-s
XX beta-s
XX bone m
XX bone m
XX bone m
XX bone m
XX Constructed by WO2001
XX WO2001
XX Constructed by WO2001
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX WO2001
XX WO300
PF 28-PEB
XX WPI; 28-PEB
XX WPI; 2
XX WPI; 2
XX New ad
PT heart
PT heart
PT partic
XX New ad
PT partic
XX The pr
CC are calls
RESULT 376
AAH44377/c
ID AAH443
XX
AC AAH443
XX
DT 26-SEP
XX
DB Beta-8
XX
W Differ
XW Differ
XW prolif
XW prolif
XW bone m
XX
XX
XX
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to cells isolated from bone marrow, which are capable of at least differentiating into heart muscle cells. The cells are applicable as remedies for various heart diseases particularly with damaged heart muscle accompanying degeneration. The present sequence is a PCR primer, which was used to illustrate the present invention
                    Differentiation; heart muscle cell; cyrokine; transcription factor; proliferation; surface antigen; heart disease; cardiomyocyte; bone marrow; umbilical blood cell; heart muscle degeneration; myocardial infarction; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New adult bone marrow-originated cells capable of differentiating interpretations heart muscle cells, applicable as remedies for various heart diseases particularly with damaged heart muscle accompanying degeneration.
                                                                                                                                                                           26-SEP-2001
                                                                                                                                                                                                                     AAH44377;
                                                                                                                                                                                                                                                              AAH44377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (KYOW ) KYOWA HAKKO KOGYO KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1185
                                                                                                                                                                                                                                                                                                                                                                                                              1185 AAGAGAGAGAGAAATCAGAG 1206
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19; Conserv
                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24 BP; 3 A; 7 C; 4 G; 10 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 146; 158pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hata J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                PCR primer SEQ ID NO:43.
                                                                                                                                                                           entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cell; heart disease; beta-skeletal actin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cardiant; cell differentiating agent,
                                                                                                                                                                                                                                                                ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.2; DB Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
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RESULT 377
ABZ55248/C
ID ABZ252
XX ABZ252
XX ABZ252
XX ABZ252
XX ABZ252
XX Human
XX Human
XX Human;
XW Human;
XW Anti-H
XX Amo anti-H
XX Amo a
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                                                                                                                                                                                                                                                                                                                                                                                                                          CC cardiomycotytes. Also described are: (1) cardiomycotytes produced by the CC differentiation of the cells; (2) a method for carrying out the CC differentiation into cardiomycotytes, regulated by a promotional and/or CC inhibitory factor; (3) a method for the differentiation of the cells into CC cell types other than cardiomycotytes; (4) drug compositions promoting the CC formation of heart muscle and regeneration of heart tissue which contain the cells; (5) a method for the production of antibodies which recognise cells; (6) a method for screening factors which promote the proliferation of the cells; (6) a method for immortalising the cells by expressing CC telomerase in them; (8) drug compositions for the treatment of heart CC disease which contain the immortalising the cells by expressing CC differentiation into cardiomyocytes. The cells are used in the treatment CC differentiation into cardiomyocytes. The cells are used in the treatment CC differentiation and in the study of cardiomyocyte differentiation. ANH44351 to AAH44409 and AAB99915 to AAB99935 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                          Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                             Human; peroxidase 9.90; enzyme; cancer;
anti-HIV; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cells capable of differentiating into cardiomyocytes and originating bone marrow or umbilical blood cells for study of cardiomyocyte differentiation and treatment of heart disease.
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Synthetic.
                                 Homo sapiens
                                                                                                              Human
                                                                                                                                               24-APR-2003
                                                                                                                                                                               ABZ25248
                                                                                                                                                                                                             ABZ25248 standard; DNA; 24 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention describes cells originating in bone marrow umbilical blood cells which are capable of differentiating into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yamada
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-FEB-2000; 2000WO-JP001148
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-NOV-2000; 2000WO-JP007741.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200148150-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (КУОМ ) КУОМА НАККО КОСУО КК
                                                                                                                                                                                                                                                                                                                          1185 AAGAGAGAGAGAAATCAGAG 1206
                                                                                                                                                                                                                                                                                                                                                        19;
                                                                                                              peroxidase 9.90 PCR primer
                                                                                                                                                                                                                                                                                            22
                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 150; 187pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hata J,
                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99JP-00372826
                                                                                                                                                                                                                                                                                                                                                                       86.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Fukuda K,
                                                                                                                                                                                                                                                                                                                                                                                        0.3%;
                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                        Score 17.2;
                                                                                                                                                                                                                                                                                                                                                                           Pred.
                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ogawa S,
                                                                                                                #2.
                                                                                                                                                                                                                                                                                                                                                                           No.
                                                                                HIV infection; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sakurada K,
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                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             မှု
                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 378
ABA01048/c
ID ABA010
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                 Human sodium pump subunit 12 of sodium pump and encoded polynucleotide, used in diagnosis and treatment of malignant tumors, hemopathy, human immunodeficiency virus infection, immunological diseases and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to human peroxidase 9.90 (see ABP5911 The peroxidase is useful for treating diseases such as cancer and H infection. The present sequence is a PCR primer, which was used in example from the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antiinflammatory; haemostatic; gene human immunodeficiency virus; HIV; inflammatory disorder; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; sodium pump; subunit 12; cytostatic; virucide; immunomodulator; antiinflammatory; haemostatic; gene therapy; cancer; haemopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human sodium pump subunit 12 PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 24 BP; 7 A; 5 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JUL-2002
                                                                                                                                                                                                                                              WPI; 2002-017442/02.
                                                                                                                                                                                                                                                                                                    Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                               27-MAR-2000; 2000CN-00115156.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-MAR-2001; 2001WO-CN000422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200177162-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA01048;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABA01048 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Polypeptide-human peroxidase protein 9.90 and polynucleotide for coding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-733654/80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-OCT-2001
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                                                                                                                                                                                                                                                                                                                                                            (SHAN-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3588 CCATGTTGCTCAGGCTAATCTC 3609
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22 CCATATTGCTCAGGCTGGTCTC 1
                                                                                                                                                                                                                                                                                                       Xie
                                                                                                                                                                                                                                                                                                                                                            SHANGHAI BIOWINDOW GENE DEV INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Page 16 (Disclosure); 31pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Xie
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BODE GENE DEV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CO LTD SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ₽P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17.2; DB 1; Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                infection; immunological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (see ABP59112) cancer and HIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              an
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Example 2; Page 16; 35pp; Chinese.

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RESULT 379
ABN66902/c
ID ABN669
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                                                                                                              Query Match
Best Local S
Matches 19
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Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             subunit 12 comprising a 110 residue amino acid sequence, fully defined in the specification, or its fragment, analogue or derivative. The polypeptide is used in the diagnosis and treatment of malignant tumours, haemopathy, human immunodeficiency virus (HIV) infection, immunological diseases and various inflammatory disorders. The present sequence is a primer used to isolate a polynucleotide encoding the polypeptide of the
                                                                                                                                                                                                                                                                                                                                             The present invention describes human macroprotein 21.78 (I). Also described is a process for preparing (I) using DNA recombination techniques. (I) and the polynucleotide sequence encoding it (II) can used in the treatment of diseases such as embryo development ____
                                                                                                                                                                                                                               Sequence 24 BP; 0 A; 1 C; 2 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New polypeptide-human macroprotein 21.78 and polynucleotide encoding it, for treating diseases such as embryo development teratogenesis and tumor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JUN-2000; 2000CN-00116981.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JUN-2000; 2000CN-00116981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CN1331245-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; macroprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human macroprotein 21.78 PCR primer 2 SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABN86902;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABN86902 standard; DNA; 24 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 24 BP; 0 A; 0 C; 4 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 19 (Disclosure); 35pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-292882/34.
                                                                                                                                                                                                                                                                                         teratogenesis and for (I), which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (BODE-) BODE GENE DEV CO LTD SHANGHAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5393 AAAAAAATACAAAAAAAGAAAAA 5414
                                                    5393 AAAAAAATACAAAAAAGAAAAA 5414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23
22
                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Xie Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAAAAAAAACAAAAAAAAACAAA 2
AAAAAAAACAAAAACAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.3%;
milarity 86.4%;
Conservative (
                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                                                                                                                                      tumours. The present sequence represents a PCR used in an example from the present invention
                                                                                                                                         0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21.78; embryo development teratogenesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>.</u>
                                                                                                              .
                                                                                                                                      Score 17.2; DB Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 17.2; DB Pred. No. 5e+02;
                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                     DB 1; Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
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                                                                                                                 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tumour;
                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                     primer
                                                                                                                                                                                                                                                                                                                                                                                  be
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RESULT 380 ABQ09325/c

27-AUG-2001; 2001WO-US026519.

28-FEB-2002

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ABQ02710/c
ID ABQ027
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AC ABQ027
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DT 11-JUN
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                                                                                                                                                                                                                                                     RESULT 381
                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to an oligonucleotide array (I) comprising at least 25 different addresses (adapter sequences) with each comprising a different capture probe selected from a group consisting of the sequences given in ABQ00010-ABQ13409. (I) is useful for immobilising a target nucleic acid sequence by attaching a adapter nucleic acid (ABQ00010-ABQ13409) to a target nucleic acid to form a modified target nucleic acid and contacting the modified target nucleic acid with (I). The steps of above method is useful for detecting a target nucleic acid, which further comprises detecting the presence of the modified target nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                       Oligonucleotide adapter/capture probe 2701.
                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 24 BP; 6 A; 9 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Array comprising adapter sequences useful for immobilizing or target nucleic acid sequence, has different addresses comprisi different specific capture probes.
                                                       WO200216649-A2.
                                                                                 Synthetic
                                                                                                             Oligonucleotide
                                                                                                                                                                   11-JUN-2002
                                                                                                                                                                                               ABQ02710
                                                                                                                                                                                                                           ABQ02710 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; Page 204; 261pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gunderson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-AUG-2000; 2000US-0227948P.
29-AUG-2000; 2000US-0228854P.
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nilarity 86.4%;
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                                                                                                                                        Array comprising adapter sequences useful for immobilizing or detecting target nucleic acid sequence, has different addresses comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
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                                                           Claim 1; Page 204; 261pp; English
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29-AUG-2000;
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29-AUG-2000;
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                                                                                                            specific capture probes
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ilarity 86.4%;
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                                                                                                                                                                                                             Sequence 24
                                                                                                                                                                                                                                                                                              The present invention provides the protein and coding sequences of human phosphatidylinositol-3 (PTDINS-3) kinase 35. The sequences can be used in the treatment of cancer, haemopathy, HIV infection, development disorders, immunological diseases and inflammation. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New human phosphatidylinositol-3 (PTDINS3) kinase 35 for diagnosing treating malignant tumor, hemopathy, human immunodeficiency virus infection, immunological diseases and various inflammations.
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                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 12; 34pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-025836/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; phosphatidylinositol-3 kinase 35; PTDINS-3 kinase 35; cancer; haemopathy; development disorder; HIV infection; immunological disease;
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                                               ТТААЛАЛАТАСАЛАЛАСАЛ 5412
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                                                                                                                                                                                                                BP; 3 A; 0 C; 1 G; 20 T; 0 U; 0 Other;
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                                                                                                                              0.3%;
                                                                                                                                                                                                                                                                     the coding
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Pred. No. 5e+02;
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                                                                                                          Gaps
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RESULT 385
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                                                                                                                                                                                                                                                                                  mucleotide sequence that encodes a Hong Kong New Gene (HKNG) I gene cassociated with bipolar affective disorder, BAD. Also included are an expression vector comprising the nucleic acid, a host cell expressing the nucleic acid, a host cell expressing the nucleic acid, an anti-HKNG1 antibody, a method of identifying modulators of HKNG1, and identifying an individual (at risk of) having HKNG1. The presence of a polymorphism that correlates with an HKNG1 allele associated with the disorder, where the presence of the polymorphism indicates that the individual (as at risk of) having HKNG1-mediated disorder. A (small) compound which modulates (inhibits or potentiates) expression of a HKNG1 gene or gene product in a human individual is useful for the treatment of a HKNG1-mediated disorder affective disorder (BAD), severe bipolar affective (mood) disorder (BP-I) and schizophrenia.
                                                                                                                                                                                                          Query Match
Best Local (
                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                       Sequence 24 BP; 8 A; 1 C; 10 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New nucleic acid molecule Hong Kong New Gene 1 (HKNG1), useful for screening for molecules which modulate HKNG1 expression for the treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rattus sp. Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HKNG1; 88; chromosome 18p; bipolar affective disorder; BAD; PC severe bipolar affective (mood) disorder; BP-I; schizophrenia; Hong Kong new gene 1; antimanic; antidepressant; neuroleptic.
               ABK86169
                                             ABK86169 standard; DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to an isolated nucleic acid molecule comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 19; Page 174; 367pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    of bipolar disorder and schizophrenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-195962/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                02-AUG-2000; 2000US-00631275.
28-NOV-2000; 2000US-00722544.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABK43308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-AUG-2001; 2001WO-US024417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200210366-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rat KHNG1 degenerate PCR primer #6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABK43308 standard; DNA; 24 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (MILL-) MILLENNIUM PHARM INC. (REGC ) UNIV CALIFORNIA.
                                                                                                                                                          1191 GAGAGAGAAATCAGAGAAAGGCAG 1214
                                                                                                                                                                                          19;
                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Freimer NB,
                                                                                                                                                                                         Conservative
                                                                                                                                                                                                          79.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novak T
                                                                                                                                                                                                                         0.3%;
                                               ВP
                                                                                                                                                                                          1; Mismatches
                                                                                                                                                                                                          Score 17.2;
Pred. No. 5e
                                                                                                                                                                                                                         DB 1; Length 24;
                                                                                                                                                                                          4
                                                                                                                                                                                          Indels
                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR; primer;
                                                                                                                                                                                          Gaps
                                                                                                                                                                                          0
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RESULT 386
ABK86168/c
ID ABK861
XX
AC ABK861
XX
DT 24-SEP
XX

standard;

DNA; 24

ВP

0

24-SEP-2002 ABK86168; ABK86168

(first entry)

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                                  S
                                                                                                                                                                                                                 CC of gene expression in one of more samples, comprising an identimer having CC a specific oligo-dT primer sequence, where the identimer comprises a CC detectable marker at its 5' end. The system is useful for identifying any CC or all genes expressed in a given in vivo or in vitro RNA sample, as well cas the relative differences in mRNA between 2 or more samples, where CC desired, for supporting discovery of new genes, and for identifying mRNA that are expressed at different levels between 2 or more samples. The new system or method addresses limitations of prior methods by comprising CC compositions and systems that incorporate new strategies where molecular or biochemical assay compositions and systems are linked to DNA or RNA CC sequence databases for optimal resource efficiency in assaying gene expression. The system has the following advantages over existing CC methods: (a) prior sequence information or clone library construction is not needed to enable the assay; (b) provides immediate sequence in formation concerning changes or differences concerning changes or differences in one assay; (c) generates cDNA fragments from all mRNA identification in and concerning changes or differences concerning changes or differences in one assay; (c) generates cDNA fragments from all mRNAs present in the complete for subsequent investigation by common molecular biology componisms lacking significant genomic sequence in formation. The present concerning can be employed in corporation.
                                                                         Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                               Sequence 24 BP; 20 A; 0 C; 1 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 11; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying and characterizing gene expression in samples, for identifying mRNAs expressed at different levels, comprises employing an identimer having a oligo-dT primer of a specific sequence and a detectable marker at its 5' end.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-508123/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-2000; 2000US-0244933P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-2001; 2001WO-US045401.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200236828-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligo dT primer #2 used in method to study gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-SEP-2002
                                                                                                                                                                                                      invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to systems for identification and characterisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GENO-) GENOMIC SOLUTIONS INC.
                                      5391 TTANAANATACAAAAAAGAAA 5412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ₹
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               dT primer; gene expression analysis; primer; ss
ω
TTAAAAAAAAAAAAAAAAA 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dombkowski AA,
                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                 0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nagel
                                                                             0; Mismatches
                                                                                                 Score 17.2; DB 1; Length 24; Pred. No. 5e+02;
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Oligo dT

primer

#1 used in method to study gene expression.

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RESULT 387
ADB92755
ID ADB927
XX
AC ADB927
XX
AC ADB927
XX
DT 04-DEC
XX
DE Human
XX
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  desired, for supporting discovery of new genes, and for identifying mRNAs that are expressed at different levels between 2 or more samples. The new system or method addresses limitations of prior methods by comprising compositions and systems that incorporate new strategies where molecular or biochemical assay compositions and systems are linked to DNA or RNA sequence databases for optimal resource efficiency in assaying gene expression. The system has the following advantages over existing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        methods: (a) prior sequence information or clone library construction is not needed to enable the assay; (b) provides immediate sequence information in addition to information concerning changes or differences in mRNA level, to determine mRNA expression level and mRNA identification in one assay; (c) generates cDNA fragments from all mRNAs present in the sample for subsequent investigation by common molecular biology techniques; and (d) does not require prior knowledge of the sequence of the genome of the organism under investigation and can be employed in organisms lacking significant genomic sequence in formation. The present sequence represents an oligo dT primer used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of gene expression in one or more samples, comprising an identimer having a specific oligo-dr primer sequence, where the identimer comprises a detectable marker at its 5' end. The system is useful for identifying any or all genes expressed in a given in vivo or in vitro RNA sample, as well as the relative differences in mRNA between 2 or more samples, where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 11; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Identifying and characterizing gene expression in samples, for identifying mRNAs expressed at different levels, comprises empidentimer having a oligo-dT primer of a specific sequence and detectable marker at its 5' end.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-2001; 2001WO-US045401.
                                   Human hereditary haemochromatosis SNP H63D
                                                                                                04-DEC-2003
                                                                                                                                                                                                                   ADB92755 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 24 BP; 3 A; 1 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to systems for identification and characterisation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-508123/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-2000; 2000US-0244933P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200236828-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  фŢ
                                                                                                                                                                                                                                                                                                                                                                             22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                 ТТАЛАЛАЛАТАСАЛАЛАДАЛА 5412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               primer;
                                                                                                                                                                                                                                                                                                                                                                             TTAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene
                                                                                                                                                                                                                   DNA; 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nagel AC;
                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 17.2; DB:
Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               analysis; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 24;
                                   probe #14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
   RESULT 388
ADG17657/c
ID ADG176
XX ADG176
XX ADG176
XX DG176
XX DG176
XX ADG176
XX ADG
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묽 S

differentiation; osteoclast precursor cell; Toll-like receptor ligand; TLR ligand; osteopathic; gene therapy; bone loss; bacterial infection; PCR; primer; ss; RT-PCR; reverse transcription PCR; mouse; murine.

Mouse toll-like receptor (TLR) 9 RT-PCR primer SeqID19.

26-FEB-2004

(first entry

ADG17657 standard;

DNA;

24

Matches

19;

Conservative

<u>.</u>.

Mismatches

ω •-

Indels

0

Gaps

0

2282 GTCCAGATGACCTCAGGAAGCA 2303

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The invention relates to a method of determination of a copy number of a copy number of an assignment of a dosage region in a sample. The methods are useful for determining the copy number of a dosage region in a sample and genotyping a target copy number of a dosage region in a sample and genotyping a target copy number of a dosage region in a sample and genotyping disease profiling, chemopredictive testing, disease profiling, compared to the compared to the copy of th
Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; human; probe; copy number determination; dosage region; risk profiling; chemopredictive testing; disease profiling; pharmacogenetic testing; genetic mutation; genetic disease.
                                                                                  Sequence 24 BP; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Determination a copy number of a dosage region, useful for disease profiling, comprises hybridizing dosage region to cross linkable probe mixture having dosage reporter probe of cross linking agent, detectable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US2003124547-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; Page 15; 22pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-709175/67.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-SEP-1998;
03-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-MAR-2002; 2002US-00093626.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peoples R, Atta RV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (PEOP/) PEOPLES (ATTA/) ATTA R '
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  label and sequence.
  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ۲× ۶
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99US-00390124.
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                                                                                  A; 5 C; 5 G; 3 T; 0 U; 1 Other;
86.4%;
     Score 17.2; DB 1; Length 24; Pred. No. 5e+02;
  5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          useful for
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ARSSULT 389
ADG75923/c
ID ADG759
XX ADG759
XX ADG759
XX Inmuno
XX Inmuno
XX Bs; no
KW prolife
XW prolife
XW renal
XX renal
XX Synthe
XX 
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to a novel method of inhibiting differentiation of an osteoclast precursor cell comprises contacting the cell with at least one Toll-like receptor (TLR) ligand, where the TLR ligand stimulates at least one TLR on the osteoclast precursor cell, thus, inhibiting differentiation of the osteoclast precursor cell. The invention may be useful for the development of compounds with an osteopathic activity or for gene therapy. The composition and methods are useful in inhibiting differentiation of an osteoclast precursor cell which may be used in the development of therapies to treat patients suffering from bone loss associated with bacterial infection and bone loss resulting from other diseases. These may also be used in identifying a Toll-like receptor that inhibits differentiation of an osteoclast precursor cell. The present sequence is that of a mouse toll-like receptor (TLR) RT-PCR primer which was used in the exemplification of the invention.
                                                                                                                                                                                                                                                                 ss; non-CpG; immunostimulatory; non-palindromic; immune response;
proliferation; differentiation; cytokine; antibody production; B-cell;
plasmacytoid dendritic cell; immunomodulator; gene therapy;
chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inhibiting differentiation of an osteoclast precursor cell, useful for treating bone diseases, comprises contacting the cell with a Toll-like receptor (TLR) ligand that stimulates a TLR on the osteoclast precursor
                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG75923;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADG75923 standard; DNA; 24 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example, SEQ ID NO 19; 72pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2003-903937/82.
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14-APR-2003; 2003US-0462859P.
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                                                                                                     WO2003101375-A2
                                                                                                                                                                                                                                                                                                                                                                                                                            Immunostimulatory non-CpG oligonucleotide IMT 178 SeqID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYPE-) UNIV PENNSYLVANIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-MAY-2003; 2003WO-US014946.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2003094857-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3127 GAGCTGGACCTGAGCTTCAATG 3148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19;
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ilarity 86.4%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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Pred. No. 5e
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RESULT 390
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XX Gene t
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XX DS 109-AUG
XX 26-FEB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention relates to novel immunostimulatory oligonucleotides that CC contain a non-palindromic sequence motif. Specifically, it refers to DNA CC oligonucleotides (without a CQG motif), which can stimulate an immune CC response in animals of the order of primate, including humans. The immune CC and antibody production in B-cells, as well as cell differentiation, cytokine CC and antibody production in D-cells, as well as cell differentiation and CC cytokine production in plasmacytoid dendritic cells. The present CC invention describes immunomodulator compositions that also comprise an CC antigen selected from, for example, viruses, bacteria, parasites, tumour CC cells and glycolipide. As such, these DNA oligos can be used in gene CC therapy for inducing B-cell activation, treating, preventing or CC ameliorating an immune system disorder or a tumoural desease including CC carcinoma. This oligonucleotide sequence is an immunostimulatory non-CpG variant DNA oligo, used in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
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                                                                                                                                                                                                                                                                      gene typing; polymorphic microsatellite loci; PML; disease predisposition; microsatellite marker; prion disease; cystic fibrosis; malignant hyperthermia syndrome; metabolic dimilk protein; hormone; transcription factor; pT7-blue-vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New immunostimulatory oligonuclectide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.
   Geldermann H,
                                                                         09-AUG-2002; 2002DE-01036711.
                                                                                                          09-AUG-2002; 2002DE-01036711.
                                                                                                                                                26-FEB-2004
                                                                                                                                                                                 DE10236711-A1
                                                                                                                                                                                                                                                                                                                                                              Cow prion protein microsatellite locus primer #90.
                                                                                                                                                                                                                                                                                                                                                                                                   29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADO81078 standard;
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                                       (UYHO-) UNIV HOHENHEIM.
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     WPI; 1996-117045/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
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01-NOV-1994;
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KAKEN PHARM
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                                                      Shimada J,
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94JP-00268603.
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                                                         Mizushima
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RESULT 392
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                                                                         Sequence 17
                                                                                                                                               The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EARA/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoletin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 17
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                                                                                                                                                                                                                                                                                                                                                                      Claim 18;
                                                                                                                                                                                                                                                                                                                                                                                                                  Enzymatic and antisense nucleic acid inhibition of repressor genes, useful for producing e.g. granulocyte colony stimulating factor protein, interferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF05470;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                               interferon alpha
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                                                                              BP; 2 A; 2 C; 0 G; 13 T; 0 U; 0 Other;
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  17;
. No.
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.5.1e+02;
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  DB 1; I
5.1e+02;
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밁
                                                                                                                                                   CC in the phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses. The invention may be useful for the CC development of compounds with a cytostatic, virucide, neuroprotective, CC probes and primers for detecting, indentifying, quantifying and/or amplifying nucleic acid, for example as one component of a gene chip, in CC witro as antisense reagents and for production of recombinant CC polypeptides. The invention may therefore be useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that CC are characterised by development of tumours or cell degeneration. The greent sequence is that of a nucleic acid sequence of the invention. CC Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO CC at ftp.wipo.int/pub/publishedpct_sequences
                                                           Query Match
Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2003025177-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human tumour suppression/reversion-related DNA sequence SeqID3660
                                                                                                                          Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; SEQ ID NO 3660; 30pp; French
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                                                                                                                                                                                                                                                                                                                                                                                                    This invention relates to novel isolated nucleic acid sequences involved
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                             5238 GATCTACAAGAGCAAGC 5254
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                                                           1 Similarity
17, Conserv
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                                                             Conservative
                                                                                                                          BP; 7 A; 4 C; 4 G; 2 T; 0 U; 0 Other;
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                                                                          0.3%;
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17
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                                                             Score 17; DB 1; Length 17; Pred. No. 5.1e+02; O; Mismatches O; Indels
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RESULT 394

Mus musculus

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AAT12446/c
ID AAT124
XX
AC AAT124
XX
XX AAT124
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AT14vi
XX AATtivi
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XX AAT124
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XX WOO9603
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CC AAT1
RESULT 395
ABA33055
ID ABA330
XX ABA330
XX ABA330
XY 11-APR
XX Mouse
XX Mouse
XX netrin
XX netrin
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Best Local Similarity
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netrin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAT12435-T12454 represent phosphorothioate oligonucleotides with antiviral activity. These sequences, and the phosphorothioate oligonucleotides represented by AAT12418-T12434 (which are complementary to regions of the mRNA 4 or 5 of herpes simplex virus 1 (HSV)), are effective in the prevention and treatment of viral infection. The sequences are especially effective against infection by HSV, HIV or varicella zoster virus (VZV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antiviral; phosphorothioate; mRNA 4; mRNA 5; herpes simplex virus 1; viral infection; HIV; varicella zoster virus; VZV; therapy; ss.
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                                                                                                                                                                                      11-APR-2002
                                                                                                                                                                                                                                       ABA93055
                                                                                                                                                                                                                                                                                        ABA93055 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 6; Page 150; 163pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antiviral phosphoro:thioate oligo:nucleotide(s) - active against e.g. herpes simplex virus 1, HIV and varicella zoster virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (LTTL-) LTT INST CO (KAKE) KAKEN PHARM
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01-NOV-1994;
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                                                                                                                             Mouse membrane bound type netrin PCR primer SEQ ID NO:20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            228 CCCTCACCCTCACCCTC 244
                                                                                                                                                                                                                                                                                                                                                                                                                           18 CCCTCACCCTCACCCTC 2
                                                                            membrane bound type netrin; glycosylphosphatidylinositol; GPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17;
                                                   B1; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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94JP-00268603.
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Pred. No.
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5.1e+02;
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RESULT 396
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XX ADC984
XX ONFK202
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                                                                                                                                                                                                                                     20-DEC-2001;
04-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NFK202 polymorphism marker PCR primer B primer seq.
                                                       WPI; 2003-559156/52
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single nucleotide polymorp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer for a mouse membrane bound type netrin invention
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                                                                                                                                                                                     INCYTE GENOMICS
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                                                                                                     Valdes A,
Schafer A;
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2002US-0423559P.
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                                                                                                                               Townley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BMD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 17;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bone damage; polymorphism; osteoporosis; SNP; PCR primer; ss; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                               Mangion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1; Le
5.2e+02;
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                                                                                                                               Galwey
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Determining whether an

individual is predisposed to

susceptibility to

low

The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions,

first active agent comprising an initiation codon, coding region,

Claim 15;

SEQ ID

NO 2468; 872pp;

English

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or

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RESULT 397
ABZ87226/c
ID ABZ87226/c
ID ABZ872
XX ABZ872
XX ABZ872
XX Human
XX Homo &
XX Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a method of determining whether an individual is predisposed to susceptibility to low bone mineral density (BMD) and/or bone damage comprising identifying whether the individual has at least one polymorphism in a polymucleotide encoding a protein, where the polymorphism in a polymucleotide encoding a protein, see ADC98235 to ADC98315). An agent identified in an method from the present invention which can be used for the prevention or treatment of a disease resulting in susceptibility to low BMD and/or bone damage is useful in the manufacture of a medicament for use in modulating the susceptibility to low BMD and/or bone damage. The disease associated with low BMD and/or bone damage is osteoporosis. The present pcR primer sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                           Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24-APR-2001; 2001US-0286137P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-APR-2002; 2002WO-US013135
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polymorphisms in associated genes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EPIG-)
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                                                                                                                                                                                                                                                                                                                   2003-229219/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17
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                                                                                                                                                                                                                                                                                                                                                               Li Y, Sa
Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 4 A; 4 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                  Sandrasagra
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  respiratory
                                                                                                                                                                                                                                                                                                                                                                        Shahabuddin
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             INC.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                  'n
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Le 5.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                     Pabalan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 20
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                                                                                                                                                                                                                                                                                                                                                                                                     Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy;
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RESULT 398
ABD23456/c
ID ABD234
XX BBD234
XX B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                          Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antisathmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                            Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic soids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-APR-2002; 2002WO-US013143.
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                                                                                                                                                 bronchodilating
                                                                                                                                                                                                                                                                                                                                     WPI; 2003-093058/08.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pulmonary transplantation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1180 AGAGAAAGAGAGAGAGA 1196
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                          ŗ
                                                                                                                                                 agent.
                                                                                                                                                                                                                                                                                                                                                                                                      Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A; 8 C; 0 G; 12 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rejection; ss; primer.
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 20; 5.2e+02;
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This invention describes a novel composition (a) a first active agent,

Differential display method for identifying secreted or transmembrane protein, comprises contacting a DNA with a first primer that hybridizes

Claim 15; SEQ ID NO 2468; 763pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 399
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Best Local
                                                                                                                                                                                                                                                                                                                                                                  Differential display method; leucine-rich motif; transmembrane protein; secreted protein; secreted protein signal peptide; ss.
                                                     P-PSDB;
                                                                                                                                                                                23-JAN-2001; 2001US-0263158P.
                                                                                                                                                                                                                    23-JAN-2002; 2002WO-IL000071.
                                                                                                                                                                                                                                                         01-AUG-2002
                                                                                                                                                                                                                                                                                                                                  Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                        DNA encoding secreted protein signal peptide sequence #13
                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-JAN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABX03804 standard; cDNA; 21
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                                                                                                                                                                                                                                                                                             WO200259259-A2
                                                                                                                                               (UYRA-) UNIV RAMOT APPLIED RES &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1180 AGAGAAAGAGAGAGAGA 1196
                                                                      2002-599769/64.
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                                                     ABG98333.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 0 A; 8 C; 0 G; 12 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 17; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                               GNI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5.2e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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RESULT 400
AAV85582
ID AAV855
XX AAV855
XX LRP5
AC LRP5
CH LRP5 P
XX LRP5;
KW INSULI
KW Glomez
KW PCR PZ
XX SYNTHE
OS SYNTHE
OS SYNTHE
OS SYNTHE
OS HOMO E
XX WO9846
XX HOS-JUN
XX (WELL
PA (WELL
PA (MERI
XX TOOM
PI HEY P
PI TOOM
PI NEW 11
XX NEW 11
XX NEW 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PEXTRACTOR OF THE STATE OF THE 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a differential display comprising contacting CC cDNA with a first primer that hybridises to an oligonucleic sequence coding for a leucine-rich motif, and with a second oligonucleotide primer to form a cDNA-hybrid molecule. The method comprises obtaining mRNA from CC entacting the cDNA with a first primer that hybridises to an CC oligonucleic sequence coding for a leucine-rich motif, and with a second cligonucleotide primer to form cDNA-hybrid molecules, amplifying the cDNA cC hybrid molecules, detecting amplified products and comparing the cDNA cC hybrid molecules, detecting amplified products and comparing the cDNA cc maplified products from each sample to identify distinctive amplified comparing the cDNA comparing the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LRP5; LDL-receptor related protein; LRP-3; IDDM; diagnosis; endocytosis; insulin dependent diabetes mellitus; autoimmune disease; glomerulonephritis; inflammation; viral infection; osteoporosis; hypercholesterolemia; Alzheimer's disease; low density lipoprotein;
                                                                                                                                                                            Todd JA, Hess JW, (Hey P, Kawaguchi Y, Phillips MS, Twells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LRP5 PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAV85582;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV85582 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       to a sequence coding for a leucine-rich motif and with a second oligonucleotide primer.
New isolated LDL-receptor related protein - used to de treating, e.g. elevated triglyceride levels, diabetes,
                                                                                                             WPI; 1998-594573/50
                                                                                                                                                                                                                                                                                                                       (WELL )
                                                                                                                                                                                                                                                                                                                                                                                                                             15-APR-1997;
05-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-OCT-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9846743-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Fig 2; 37pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2639
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 CCCTGCAGCTGCTGCTG 18
                                                                                                                                                                                                                                                                                                                       WELLCOME TRUST LTD.
MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 100.
17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCCTGCAGCTGCTGCTG 2655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21 BP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0043553P.
97US-0048740P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 7 C; 8 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                Caskey CT,
Y, Merriman
                                                                                                                                                                               RCJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
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Pred. No.
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                                                                                                                                                                                                                RD, Gerhol
Metzker ML,
                                                                                                                                                                                       Gerhold D, наш...
чт. Nakagawa y
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .3e+02;
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                                develop products
                                                                                                                                                                                                                                                     Hammond
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The present invention describes IRP5 (low density lipoprotein (LDL) receptor related protein, previously designated LRP-3). AAV85525 to AAV85586 represent PCR primer used for obtaining LRP5 cDNA. Nucleic acid molecules (NAMS) encoding LRP5 can be used for determining if an individual is susceptible to insulin dependent diabetes mellitus (IDDM). The NAMS or proteins can be used for reducing triglyceride levels in the serum of an individual. Therapies that affect LRP5 may also be useful in the treatment of autoimmune diseases such as glomerulonephritis, diseases and disorders involving disruption of endocytosis and/or antigen presentation, cytokine clearance and/or inflammation, viral infection, pathogenic bacterial toxin contamination, elevation of free fatty acids or hypercholesterolemia, type 2 diabetes, osteoporosis, Alzheimer's disease and cardiovascular disease. Products from the present invention
Sequence
                                                        can also
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 12; Page 98; 200pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disorders, inflammation or Alzheimer's disease
     20
                                                        be used for detection, diagnosis and drug screening
  BP; 4 A; 6 C; 5 G; 5 T; 0
     U; 0 Other;
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RESULT 401
AAF23284/c
ID AAF232
XX AAF232
XX ITS; 1
XW MYCODB
XW MYCODB
XW MYCODB
XW MYCODB
XX ISJMI
PR 29-MAI
PR 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide for detection of Mycobacterium smegmatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF23284;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mycobacterium
Mycobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               internal transcribed spacer region; Mycobacterium fortuitum; bacterium chelonae; Mycobacterium abscessus; Mycobacterium vaccbacterium ellavescens; Mycobacterium astaticum; tuberculosis; bacterium porcinum; Mycobacterium acapulcensis; identification; bacterium diernhoferi; PCR primer; probe; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AGGCTAATCTCAAACTCCTG 3618
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGGCTGGTCTCAAACTCCTG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
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Query Match Best Local S Matches 18

Local Similarity

0.3%; nilarity 90.0%; Conservative

0

Score 16.8; DB 1; Pred. No. 5.6e+02; Mismatches

Length Indels

20;

0

Gaps

0

3.6e+02; 2;

18;

Mycobacterium smegmatis. Mycobacterium

16-MAY-2000; 2000WO-KR000477.

29-MAY-1999; 29-MAY-1999; 29-MAY-1999; 29-MAY-1999; 99KR-00019632. 99KR-00019633. 99KR-00019634. 99KR-00019635. 99KR-00019631.

07-APR-2000; 29-MAY-1999; 2000KR-00018189

(SJHI-) KIMC/)) SJ HIGHTECH |) KIM C M.) PARK H K. ႘

LID.

Kim CM, Park HK, Jang 품

WPI; 2001-061527/07

Novel oligonucleotide sequences of internal transcribing spacer region non-tuberculosis mycobacteria (NTM) used as probes or primers for detecting and identifying mycobacteria and distinguish TB complex from

Claim

29; Page 68; 89pp; English

0

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RESULT 402
AAF75041
ID AAF750
XX AAF750
XX DE-MAY
XX S-hyda
XX S-hyda
XX Homo s
XX The s
CC Gene
CC Gene
CC Gene
CC In add
CC In add
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mycobacterium ITS (internal transcribed spacer region) nucleotide sequence. ITS DNA sequences from M. fortuitum, M. chelonae, M. abscessus, M. vaccae, M. flavescens, M. asiaticum, M. porcinum, M. acapulcensis, M. diernhoferi genes were identified. The oligonucleotides derived from these sequences were used to develop PCR primers and hybridisation probes for detection and identification of Mycobacterium. ITS has a more polymorphic region than 165 rRNA and also has a conserved region. It is therefore highly effective as a target DNA for distinction of genotype. The oligonucleotide probes, attached to solid substrate, hybridise only the therefore and the probes attached to solid substrate, hybridise only the theory of the third they have the theory of the third they have the third they have the theory of the third they have the theory of the theory of the third they have the theory of the
                       The present invention relates to 5-hydroxy tryptamine receptor 1A (HTR1A) gene. HTR1A-encoding polynucleotides containing one or more of the novel polymorphic sites are useful in studying the expression and biological function of HTR1A, as well as in developing drugs targeting this protein. In addition, information on the combinations of polymorphisms in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      with nucleotide sequences in ITS of specific mycobacteria, and thus they can detect and identify the specific mycobacteria sensitively. The oligonucleotides can also detect and identify the specific mycobacteria by PCR amplification. Using the oligonucleotide primers or probes made from ITS of mycobacteria, it is possible to detect mycobacteria, distinguish tuberculosis (TB) complex from non-tuberculosis mycobacteria (NTM), and to identify mycobacteria species accurately and effectively
                                                                                                                                                                                                                                                                                      New 5-hydroxy tryptamine receptor 1A gene variants expression and biological function of the gene and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5-hydroxy tryptamine receptor IA; HTRIA; polymorphism; Tourette's; neuropsychiatric; ss.
                                                                                                                                                                                                Example 1; Page 33; 64pp;
                                                                                                                                                                                                                                                        targeting 5-hydroxy tryptamine receptor 1A protein
                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-191514/19
                                                                                                                                                                                                                                                                                                                                                                                                                                 Denton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-AUG-2000; 2000WO-US040519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200110884-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer #13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF75041;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mycobacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-FEB-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              193 CGTTGCCCACACCCCATCTC 212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10,
                                                                                                                                                                                                                                                                                                                                                                                                                                 ₽₽,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CGTTTCCCACACCGCATCTC 1
  may have diagnostic and forensic applications. A
                                                                                                                                                                                                                                                                                                                                                                                                                                 Kliem SE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B₽;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 C; 10 G; 3 T; 0 U; 0 Other;
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Pred. No. 5.6e+02;
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                                     19-JAN-2000; 2000US-00487445
                                                                         19-JAN-2000; 2000US-00487445.
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(ISIS-) ISIS PHARM INC

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16-OCT-2000;
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16-OCT-2000;
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                          Grosse WM,
Kekuda R,
Edinger S,
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16-OCT-2000;
16-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antidiabetic; immunosuppressive; neuroprotective; gene therapy; cancer; cardiomyopathy; atherosclerosis; cell signal processing; diabetes; AIDS; metabolic pathway modulation; neoplatic; neurological disorder; asthma; adonoscriptums.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; NOVX; cytostatic; antiarteriosclerotic; cardiovascular; lymphoma; antidiabetic; immunosuppressive; neuroprotective; gene therapy; cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zhang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18-APR-2002
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Crohn's disease; multiple sclerosis; Graft versus host disease;
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                       Alsobrook JP, Lepley DM, Burgess CE, Mishra
Li L, Padigaru M, Shimkets RA, Zerhusen BD,
Gerlach V, Macdougall J, Stone D, Gunther E,
                                                                                                                                                                                                                      ; 2000US-0240113P.
2000US-0240625P.
2000US-024063PP.
2000US-0240648P.
2000US-0240669P.
2000US-0240669P.
2000US-0240732P.
2000US-0241190P.
2001US-0262455P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes novel human proteins designated NOVX (where X is 1, 2a, 2b, 2c, 2d, 3, 4, 5, 6a, 6b, 7, 8, or 9). NOVI is a C tyrostine-protein kinase 6-like protein; NOV2a-d are keratin 4-like protein; NOV3 is a collagen-like protein; NOV4 is a cystatin B-like CC protein; NOV5 is a serotonin receptor-like protein; NOV6a and NOV65v are CC cold inducible glycoprotein 30-like proteins; NOV7 is a matrilin-2-like protein; NOV8 is a leukocyte surface antigen (CD53)-like protein; and CC protein; NOV8 is a leukocyte surface antigen (CD53)-like protein; and CC entiarteriosclerotic, cardiovascular, antidiabetic, immunosuppressive and consurprotective activities, and can be used in gene therapy. The NOVX sequences can be used in therapeutics, particularly for treating, constate in a subject, particularly a human. These disorder or a pathological cardiomyopathy, atherosclerosis, a disorder related to cell signal cC processing and metabolic pathway modulation or diabetes. The NOVX sequences are also useful for determining the presence of or processing and in the rapeutic or prophylactic actions for conjugatic or neurological disorders, and in the treatment of conjugation or neurological disorders, and in the treatment of consequences are also useful in therapeutic or prophylactic applications for consequences are also useful in therapeutic or prophylactic applications for consequences are sequences, and in the treatment of conjugations, lymphoma, prostate cancer, uterus cancer, immune conjugations of the present sequences, multiple sclerosis or Graft versus host disease. The present sequence represents a PCR primer for buman NOV7. which is used in an example from the present a PCR primer for conjugations of the present and the present and invent in our and an example from the present and invent invention.
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Plant growth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cancer or diabetes.
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Pred. No. 5.6e+02;
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28-MAR-2001; 2001JP-00091994.

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(TOYT) TOYOTA JIDOSHA

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2003-170475/17.

stress-resistant plant comprises a gene encoding a protein

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                                                                                                                                                                                                                                                 The present sequence is that of sequencing primer OVMU44, which was used to sequence the chicken ovomucoid gene expression control region (see ACC84070). A claimed nucleic acid comprises the ovomucoid gene expression control region operably linked to a nucleotide sequence encoding a heterologous polypeptide, especially an immunoglobulin subunit or interferon alpha-2b. The nucleic acid is used to express the heterologous polypeptide in a host (especially avian) cell or in a transgenic avian, which produces the heterologous polypeptide in the serum or egg white
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated nucleic acid comprising avian ovomucoid gene expression control region, useful for directing expression of a nucleic acid encoding a polypeptide e.g., immunoglobulin, in an avian, preferably chicken cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to proteins with peroxidase activity and their coding sequences (ABZ25464-ABZ25465 and ABP70669-ABP70670). The peroxidase coding sequences were used to generate a transformed stress-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Harvey AJ,
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                                                                                                                          Pred.
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The invention relates to a compound 8-30 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding caspase 8 (a protein involved in apoptosis), and inhibits the expression of caspase 8, i.e. an antisense oligonucleotide. Also included are a compound 8-30 nucleobases in length that specifically hybridises with at least an 8-nucleobase portion of an active site on a nucleic acid molecule encoding caspase 8, a composition comprising the compound and a carrier or diluent, inhibiting the expression of caspase 8 in cells or tissues (by contacting the cells or tissues with the compound so that expression of caspase 8 is inhibited) and treating an animal having a disease or condition associated with caspase 8 by administering to the enimal a therapeutic or prophylactic amount of the compound so that expression of caspase 8 is inhibited. The compound, composition and methods are useful for treating a disease or condition associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Caspase-8; cytostatic; immunosuppressant; anti-HIV; ss; antisense gene therapy; apoptosis; hyperproliferative disorder; haematopoletic disorder; autoimmune disorder; viral infection; AIDS; neurological disorder; Alzheimer's disease; Parkinson's disease; amyotrophic lateral sclerosis; retinitis pigmentosa; blood cell disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding caspase 8, useful for treating a disease/condition associated with caspase 8, such as hyperproliferative or autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15;
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11-JAN-2001; 2001WO-US000955
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(COWS/)
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COWSERT L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cowsert LM;
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The invention relates to new antisense compounds targeted to a nucleic acid molecule encoding human cholesteryl ester transfer protein, specifically hybridises with it and inhibits the expression of human cholesteryl ester transfer protein. The compound is useful for preparing a composition for treating abnormal lipid or cholesterol metabolism, atherosclerosis or cardiovascular disease. The present sequence
                                                                                                                                                                                                                                           New antisense compound, usefu abnormal lipid or cholesterol cardiovascular disease.
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Pred. No. 5
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Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel method for establishing an expression system of spider drag-line silk genes in silkworm by fusing the silkworm fibroin L-chain cDNA and its promoter upstream of the spider drag-line silk gene cDNA, ligating the fused gene with a reporter gene and inserting into a transposon to obtain a recombinant transposon which can be used to transform a silkworm egg. The method of the invention is useful for establishing an expression system of spider drag-line silk gene in B. mori. The spider dragline silk gene product accounts for 30% of total silk proteins. This method provides a rate of transformation of about 0.5-1%. The present sequence represents a PCR primer used to amplify the silkworm spider dragline silk gene (MaSPI) sequence used in
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Silkworm; primer; 88; spider drag-line; silk; fibroin; PCR; light chain;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amplify the silkworm spider the method of the invention
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Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, ob cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                                                     antisense oligonucleotide; glucocorticoid receptor; infection;
inflammation; tumour formation; diabetes; obesity;
                                                                                                                                                                                                                                                                                                                                                        ADH65015 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                  #1849
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RESULT 412
ADK7444/c
ID ADK744
XX ADK744
XX Chimer
AC Chimer
XX Nav1.3
KW diabet
KW infant
XX Synthe
PN WO2004
XX WO2004
XX 14-AUG
PR 14-AUG
XX ROBERG
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Best Local S
Matches 18
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ئار
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention seeful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothicate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New antisense compound targeted to a nucleic acid molecule enc mammalian glucocorticoid receptor, useful for treating diabete cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                         14-AUG-2002; 2002US-0403416P
                                                                                                                                                26-FEB-2004
                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                    diabetic neuropathy; arthritic pain;
                                                                                                                                                                                                                                                                   Navl.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia;
                                                                                                                                                                                                                                                                                               Chimeric phosphorothioate oligonucleotide to target Nav1.3 #1778.
                                                                                                                                                                                                                                                                                                                              20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                         ADK74444;
                                                                                                                                                                                                                                                                                                                                                                                       ADK74444 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 1 A; 12 C; 0 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; SEQ ID NO 1849; 985pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Сговьу
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAY-2002; 2002US-0381857P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-MAY-2003; 2003WO-US016084.
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                                                                                                                   14-AUG-2003; 2003WO-US025465
                                                                                                                                                                             WO2004016754-A2
                                                                                                                                                                                                                                     infantile epilepsy; ataxia; ss.
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                                                           (PHAA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2565 GGGGGAGAGAGAGATGGAGA 2584
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                                                           PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGGGAAGAGGGAGATGGAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                        DNA;
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                                                                                                                                                                                                                                                                                                                            entry)
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.8;
No. 5
                                                                                                                                                                                                                                                      migraine headache
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          diabetes,
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WPI; 2004-203785/19

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RESULT 413
ADP11348/c
ID ADP113
XX ADP113
XX ADP113
XX Transpan
CX Transpan
XX Iranspan
XX Wollam
XX 21-MAJ
XX 21-MAJ
XX 21-MAJ
XX Wollam
XX Wollam
XX Wollam
XX WPI;
XX Diagnx
PT Pancri
PT rejecti
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Best Local S
Matches 18
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                                                                                                                                                                                pancreas, pancre
                                                                                                                                                                                                                                                       Diagnosing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24-APR-2002; 2002US-00131831.
20-DEC-2002; 2002US-00325899.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transplant rejection;
inflammatory bowel dis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Taqman probe of the invention #31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADP11348;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADP11348 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense compound targeted to a nucleic acid molecule encoding Navl.3, useful for useful for treating a disease or condition associated with Navl.3, e.g. pain, seizure disorder such as childhood seizure
                                                                                                                                                                                                                                                                                                                                  WPI; 2004-400724/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Wohlgemuth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EXPR-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EXPRESSION DIAGNOSTICS INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GAAAAGATGGCACAGGCACT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
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                                                                                                                                                                                   pancreatic
, in an indi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              or ataxia.
                                                                                                                                                                                or monitoring transplant rejection, e.g. heart, kidney, liver, pancreatic islet, lung, bone marrow or stem cell transplant in an individual, comprises detecting the expression level of
                                                                                                                                                                                                                                                                                                                                                                                                                                       Ģ
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Fry K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   on; immune system; rheumatoid arthritis;
disease; multiple sclerosis; HIV; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Woodward
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Pred. No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Prentice
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                                                                                                                                                                                                                                                                                                                                                                                                                                           Morris
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Claim 58; SEQ

ID NO 1357; 1762pp; English

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ID AAQ919
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AC AAQ919
XX T-cell
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XX T-cell
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XX Synthe
XX W09500
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XX U00500
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                                                                                       TCR Va families expressed in human uveal melanoma were analysed by PCF amplification. In 7 of the 8 cases, the Va7 region was expressed and rearranged. The identity was confirmed by hybridisation to a Va7 probe (AAQ91937). The variable regions associated with different neoproliferative tissues may be determined in this way; a homogeneous composition of T-cells may then be administered for treatment of the particular neoproliferative tissue. (Updated on 25-MAR-2003 to correct
                                                                                                                                                                                                                                                                                                                                                                                                 Determining relation between auto:immune degenerative diseases and specific variable regions of T-cell receptors - as associated with the host HLA or T-cells associated with combating neoproliferative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                T-cell receptor Va7 hybridisation probe.
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28-NOV-1995
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                                                                                                                                                                                                                                                                                                                                         Example 1; Page 35; 122pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oksenberg
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TCR; variable region Va7; multiple sclerosis; urodegeneration; uveal melanoma; ss.
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Pred. No. 5.6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mitchell
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probe
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RESULT 415
AAQ75729/c
ID AAQ75729/c
ID AAQ757
XX AAQ757
XX AAQ757
XX ATALYS
KW ATALYS
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FOR O1-NOV
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FOR WPI; 1
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CC ATALYS
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CC LIABEL!
CC CLARAC
CC CATAL
CC CAT
RESULT 416
AAT92734
ID AAT927
XX
AC AAT927
XX
DT 25-MAR
DT 04-FEB
XX
DE V-alph
XX
FODE;
KW Probe;
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Best Local Similarity
Matches 18; Conserv
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GRNESEQ files AQ75547-Q75788) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 2 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ75729 standard;
Probe; hybridise; T-cell receptor; TCR V-alpha; TCR V-beta; brain; MS;
                                                        V-alpha7 probe for T-cell receptor
                                                                                                                 25-MAR-2003
04-FEB-1998
                                                                                                                                                                                                     AAT92734
                                                                                                                                                                                                                                                              AAT92734 standard; cDNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Analysis of cDNA and gene expression - by amplification of by digestion with restriction enzymes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              04-AUG-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              5389 AATTAAAAAAATACAAAAA 5408
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                                                                                                              (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 16.8; DB 1; Length 21; Pred. No. 5.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.8; DB 1;
Pred. No. 5.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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RESULT 417
AAT76098/c
ID AAT76098 standard; D
XX
AC AAT76098;
XX
DT 12-SEP-1997 (first
XX
DE Human histidine deca
XX
KW Asthma; airway epith
XX
KW chronic obstructive
XX
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                                                                                                                                                                                                                                                                                                               ঠ
                                                                                                                                                                                                                                                                                                                                                                                                                             AAT92734 and AAT92735 represent probes for the V-alpha chain of the T-CC cell receptor (TCR). This sequence can be used in the method of the invention is for determining the presence, in a human host, of T-cells associated with multiple sclerosis (MS). The method comprises isolating T-cells from the brain or cerebrospinal fluid of a human host, and detecting in the T-cells the presence of a limited number of rearranged complementarity determining region 3 (CDR3) regions of the TCR V-alpha or V-beta chains. The rearrangements that are detected are associated with MS. The detection is carried out by isolating nucleic acid molecules from the TCR, and amplifying the molecules with primers specific for sequences 5' and 3' of the rearranged CDR3 region (such as AT92727-T92757). The method can be used for the diagnosis of MS. In CC AAT92727-T92757). The method can be used for the diagnosis of MS. In addition, by identifying specific TCR variable regions associated with the disease, therapies may be employed to inhibit the attack of the T-CC cells having such variable regions on the target cells or proteins. The therapies may involve ablation of T-cells carrying the particular variable regions, administration of compounds which inhibit binding of the T-cell receptor to the target cell, or prevention of the degenerative effects of the binding of the T-cell to the target cell or protein.

CC (Updated on 25-MAR-2003 to correct PF field.)
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Matches
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01-MAY-1991;
30-APR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Diagnosis of multiple sclerosis - by detection of T-cell receptor V-alpha or V-beta rearrangements in T-cells from the brain or cerebrospinal fluid.
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 5 A; 4 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  T-cell detection; multiple sclerosis; cerebrospinal fluid; human; CDR3; therapy; T-cell ablation; complementarity determining region 3; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example; Col 18; 52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1997-470032/43.
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                                                                                                                                                                                                                                                                                                                                                                        Local
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                                                                                                                                                                                                                                                                                                                                                                        Similarity
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                                                                                                                                                                                                                                                                                                                                                       Conservative
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91WO-US002991.
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                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                      90.0%;
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                                                                                                                                                                                                                                                                                                                                                                      Score 16.8; DB 1;
Pred. No. 5.6e+02;
                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                       DB 1; Length 21;
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Human histidine decarboxylase antisense oligonucleotide HUMHDCAS2.

(first entry)

Asthma; airway epithelium; adenosine free; cystic fibrosis; chronic obstructive pulmonary disease; bronchitis; ss.

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RESULT 418
AAV27994
ID AAV279
AC AAV279
XX ATAXIA
XX Se; PC
KW radiat
XX Synthe
OS Homo E
XX WO9822
XX WO9822
XX PD 28-MA3
XX PD 28-MA3
XX YO 28-MA3
XX YO 20-NOV
XX PA (VIRG-
XX PA (VIRG-
XX PD COncar
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A method for treating airway disease in a subject has been produced, which involves the topical administration of an essentially adenosine free antisense oligonucleotide (ON) to the airway epithelium of the subject. The present sequence is an antisense oligonucleotide HUMHDCAS2 specific for the human histidine decarboxylase. The method can be used to treat airway diseases such as cystic fibrosis, asthma, chronic obstructive pulmonary disease, bronchitis and other airway diseases characterised by an inflammatory response. By eliminating adenosine from the antisense ON, its liberation upon antisense degradation is prevented, thereby preventing adenosine- induced bronchoconstriction in patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nyce
                                                                 20-NOV-1996;
                                                                                                                    28-MAY-1998
                                                                                                                                              WO9822621-A1
                                                                                                                                                                                                            radiation;
                                                                                                                                                                                                                                                  Ataxia telangiectasia exon 19 primer 1.
                                                                                                                                                                                                                                                                                                                              AAV27994 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 0 A; 10 C; 1 G; 10
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                                       (VIRG-) VIRGINIA MASON RES CENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            with hyper-reactive airways
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYEC-) UNIV EAST CAROLINA.
                                                                                                                                                                                                                          PCR;
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                                                                                                                                                                      sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                      CAGAGAAAGAGAGAGAGA 1198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Page 26; 71pp;
                                                                                                                                                                                                                        primer; amplification; ataxia telangiectasia;
                                                                                                                                                                                                                                                                                                                                                                                               CAGAGAGAGAGAGAGAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Metzger WJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       of airway diseases such as asthma - by topically applying free antisense oligo:nucleotide to airway epithelium of
                                                                                                                                                                                                             breast
                                                                                                                                                                                                                                                                                                                                                                                                                                              0.3%;
ilarity 90.0%;
Conservative
                                                                                                                                                                                                                                                                            (first
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                                                                 96US-00753147.
                                                                                          97WO-US020953
                                                                                                                                                                                                                                                                            entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 16.8; Di
Pred. No. 5.6e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  T; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            6e+02;
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                                                                                                                                                                                                                        diagnosis;
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The primers AAV27964-V28086 are used to amplify ataxia telangiectasia (ATM) exons and their adjacent splice junction sites. These can be used as a method of detecting a mutation in the ATM gene by comparing the PCR products of amplification from a sample from a patient suspected of having an ATM mutation with a sample from a non-mutated ATM patient This method is especially useful for diagnosing ataxia telangiectasia in heterozygotes and can be used to locate the positions of the mutation. The diagnosis of ataxia telangiectasia in patients needing therapeutic radiation will prevent fatal radiation burns and the development of breast cancer which can occur
The present sequence represents a PCR primer for the Streptococcus pneumoniae phoH protein, a member of the ATP-binding protein family. The PhoH protein is administered to treat individuals in need of phoH proteins (directly or via a vector i.e. gene therapy), and as an antigen for inducing an immunological response (administered directly i.e. vaccine). They can prevent adhesion of bacteria to matrix proteins, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Streptococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 6; Page 6; 47pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Method of detecting ataxia telangiectasia - comprises use of primers based on intron-exon boundaries, useful for diagnosing disease in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-312503/27.
                                                                                                                                     New Streptococcus pneumoniae phoH polypeptide and polynucleotide - useful as diagnostic reagents and for prevention and treatment of Streptococci infections which cause conjunctivitis and meningitis.
                                                                                                                                                                                                                                                                                                                                                                                                                  Streptococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Streptococcus
ATP-binding pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX19906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX19906 standard;
                                                                                                                                                                                                                                 Zalacain M,
                                                                                                                                                                                                                                                                                              18-SEP-1997;
                                                                                                                                                                                                                                                                                                                            04-SEP-1998;
                                                                                                                                                                                                                                                                (SMIK ) SMITHKLINE BECKMAN CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3639 AATTGCTGAGATTGCAGAGG 3658
                                                                                                                                                                                                     1999-182905/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AATTGCTGAGATTACAGATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                              cus pneumoniae; phoH; antibacterial; infection; pneum
g protein; otitis media; conjunctivitis; bacteraemia;
pleural empyema; endocarditis; meningitis; PCR prime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 7 A; 2 C; 5 G; 7 T;
                                                                                                          Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                 Brown JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                   pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            pneumoniae ATP-binding phoH gene PCR primer
                                                                                                                                                                                                                                                                                            97US-00932978.
                                                                                                                                                                                                                                                                                                                          98EP-00307171.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                       25pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                          English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 16.8;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5.6e+02;
2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer; 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pneumonia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
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RESULT 420
AAX53903/c
ID AAX539
XX AAX539
XX AAX539
XX AAX539
XX DISTINCT OS-JUL
XX AAX539
XX AAT18e
KW IMPAIR
KW IMPAIR
KW PULMON
KW CCHONI
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KW PULMON
KW POSSTA
XX COLON
XW WOO913
XX WOO913
XX WOO913
XX IT-SEP
XX WPI; 1
XX WPI; 1
XX WPI; 1
XX PI VASOCC
XX DISCIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    infection. They are also useful for identifying agonists and antagonists by screening host cells expressing phoH protein, and detecting the absence or presence of phoH activity. Agonists and antagonists are useful for inhibition and treatment of conditions associated with phoH imbalance, and are therefore potential antibacterial compounds. PhoH polynucleotides are useful for genetic immunisation, preferably via a vector. Anti-phoH antibodies induced by the protein are useful for preventing or treating infections, especially bacterial infections, and also for isolating clones expressing phoH protein, or for purifying the prevent by affinity chromatography. Diseases can be diagnosed by the presence of phoH nucleic acid, and/or analysing for the presence or amount of phoH protein in a sample, due to an infection of an organism with the phoH gene. The stage and type of infection can be determined. Diseases prevented, diagnosed and treated include those caused by bacterial infection, especially Streptococci pneumoniae infection which causes otitis media, conjunctivits, pneumoniae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis; acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasoconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer;
                                                                                                                                                                                                                                                                                    17-SEP-1997;
09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                                               17-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9913886-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Histidine decarboxylase receptor antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX53903 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21 BP;
                                                                           New antisense oligonucleotides used in treatment of,
                                                                                                                                WPI; 1999-229400/19
                                                                                                                                                                                                                                                                                                                                                                                                                  25-MAR-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              05-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bacteraemia, sinusitis, pleural empyema, endocarditis and especially
                                                     vasoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hepatocellular carcinoma; kidney cancer; melanoma; hepatic metastasis;
                                                                                                                                                                                                                                    (UYEC-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               are useful for use on wounds and body implants to prevent bacterial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4932 GAACTITGATGATGCCTTTG 4951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GACCTIGGATGATGCCTTTG 2
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                                                                                                                                                                                                                                    EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                    97US-0059160P.
98US-00093972.
                                                                                                                                                                                                                                                                                                                                                               98WO-US019419
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 5.6e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>ب</u>
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                                                                           e.g. pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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Disclosure; Page 45; 120pp; English

bronchitis,

emphysema, respiratory distress

syndrome,

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RESULT 421
AAA33346/c
ID AAA333
XX AAA333
XX AAA333
XX Low ad
XX Low ad
XX Human;
KW phosph
KW allerg
KW allerg
KW lung d
KW respix
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Best Local Similarity
Matches 18; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            XAAAAAAAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The specification describes antisense oligonucleotides (AAX52869-X55271) directed against at least 2 mRNAs selected from target genes, coding and non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-771) can be used for the antisense treatment of diseases and conditions arypical diseases and conditions are those associated with impaired respiration and inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, allergic rhinitis, and continue as the antise as the continue as the conti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  acute asthma, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease (COPD), and cancers such as leukemias, lymphomas, carcinomas e.g. colon cancer, breast cancer, lung cancer, pancreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma, hepatic metastases, as well as all types of cancers which may metastasize or have metastasized to the lungs, including breast and prostate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; adenosine receptor; low adenosine antisense oligonucleotide; phosphorothicate; impaired respiration; infilammation; allergy; allergic disease; bronchoconstriction; inhibitor; antiaflammatory; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; antiallergic; antiasthmatic; cytostatic; analgesic; impaired airway; lung disease; isohaemic condition; pulmonary vasoconstriction; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD; cancer; leukaemia; lymphoma; carcinoma; metastasis; ss.
                                                                                                                                 New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension,
                                                                                                                                                                                                                                                                      WPI; 2000-205971/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200009525-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Low adenosine antisense oligonucleotide SEQ ID NO:1035
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYEC-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1179 CAGAGAAAGAGAGAGAGA 1198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             98US-0095212P
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Pred.
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No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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Claim 18; Page 394; 1343pp; English.

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RESULT 422
AAF19468/c
ID AAF194
XX AAF194
XX I AAF194
XX I AAF194
XX I Low ad
KW Low ad
KW I Low ad
KW I RESPIT
KW SURFAC
KW RESPIT
KW CARCEL
KW C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Low adenosine antisense oligonucleotide; phosphorothioate; allergy; human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary obstruction; asthma; RDS; respiratory distress syndrome; pain; cystlc fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; pulmonary transplantation rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 0 A; 10 C; 1 G; 10
                                                                                                                                                             06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chronic obstructive pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human histidine decarboxylase polynucleotide fragment #1035.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF19468 standard;
                                                                                                                                                                                                                                          24-MAR-2000; 2000WO-US008020
                                                                                                                                                                                                                                                                                                                                                                                                    WO200062736-A2
                                            (UYEC-) UNIV EAST CAROLINA.
(NYCE/) NYCE J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1179 CAGAGAAAGAGAGAGAGAGA 1198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                             99US-0127958P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90.08;
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Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   pulmonary infection; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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RESULT 423
ABL43857
ID ABL438
XX ABL438
AC ABL438
XY 11-APR
XX Human;
KW Human;
KW PCR px
XX PCR px
XX JP2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CC immunoglobulins and antibodies, antibody receptors, cytokines and CC chemokines, endogenously produced specific and non-specific enzymes, Cb binding proteins, adhesion molecules and their receptors, cytokine and CC chemokine receptors, adenosine receptors, bradykinin receptors, central nervous system (CNS) and peripheral nervous and non-nervous system peptide CC transmitters, defensins, growth factors, vasoactive peptides and CC receptors, binding proteins and malignancy associated proteins. The CC antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction and/or lung inflammation, allergy(ies) and/or Surfactant hypoproduction which are associated with a disease or CC condition selected from pulmonary vasoconstriction, inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome (RDS), pain, cystic fibrosis (CP), allergic rhinitis (AR), pulmonary transplantation rejection, pulmonary infections, bronchitis, CC and/or cancer. AAF18434 to AAF21543 represent human polynucleotide fibrosis in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                     Human chromosome 1p36-35 PCR primer SEQ ID NO:901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (I) can have respiratory, bronchodilator, antiinflammatory, analgesic, immunosuppressive, antiasthmatic, hypotensive and cytostatic activities. The antisense oligonucleotides and (I) can be used to down-regulate the expression and or activity of target polypeptides associated with lung/respiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, in the such as stimulating and activating peptide factors and transmitters.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Low adenosine (A) content antisense oligonucleotides which do not trigger adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions.
                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                    11-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                   ABL43857
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABL43857 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonuclectides and compositions (I) comprising them. In the antisense oligonuclectides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 14; Page 141; 1592pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-679539/66.
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                                                                                                                                                  JP2001321190-A.
                                                                                                                                                                                                                                                                    Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present invention describes low adenosine (A) content antisense conucleotides and compositions (I) comprising them. In the antisense
                                                                                                                                                                                                                                             primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1179 CAGAGAAAGAGAGAGAGAGA 1198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 0 A; 10 C; 1 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90.0%;
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0,

10-MAR-2000; 2000JP-00066716. 12-MAR-2001; 2001JP-00068285. 20-NOV-2001.

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RESULT 424
ABZ95162/c
ID ABZ951
XX ABZ951
XX ABZ951
XX ABZ951
XX Human
XX Homo E
X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a method of arraying genome clones. The CC method comprises: (a) clones of the genomic libraries contained in CC multiwell plates numbered for discrimination are mixed in each of the CC multiwell plates (b) a primer designed based on the chromosome marker CC sequence is added to the mixture to carry out an amplification reaction; CC (c) a signal corresponding to the marker is detected from the resultant CC amplified product to specify the discrimination Nos. of the multiwell CC plates containing the clones having said marker sequence; (d) the order CC of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell CC plates; (e) the clones in the multiwell plates of the specified CC discrimination Nos. are mixed respectively in each wells of longitudinal CC and lateral directions; (f) the mixed clones are cultured and the CC resultant cultures are amplified by using the above primer; (g) signals CC are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are cultured as the positions on the chromosome and arrayed. The CC represent pck primers for human chromosome 21422.1, which are Specifically claimed for use in the present invention
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABZ95162 standard; DNA; 21
      Мусе JW,
                                                                                                                                                                                                    23-APR-2002; 2002WO-US013135
                                                                                                                                                                                                                                                                                                                               WO200285308-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human histidine decarboxylase antisense fragment no.1025.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABZ95162;
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                                                                                                                                  24-APR-2001; 2001US-0286137P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-144136/19.
                                                                    (EPIG-) EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3600 GGCTAATCTCAAACTCCTGG 3619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inflammation; respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GGCTAATCTTGAACTCCTGG 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18;
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GENOTEX YG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
   Li Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
   Sandrasagra A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; 5 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ВÞ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
      Katz
   'n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length 21;
   Pabalan J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
   Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                        Tang L,
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Disclosure; SEQ ID NO 10404; 872pp; English.

junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject setissue, or treating bronchoconstriction, and inflammatory ensisted or stassue, or treating bronchoconstriction, and inflammatory ensisted or stassue, or treating bronchoconstriction, and inflammatory ensisted or a stassue, or treating bronchoconstriction, and the producing sensitivity to adenosine or the lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of

밁 S Matches Query Match Best Local Sequence 21 BP; 0 A; 10 C; 1 G; 1179 CAGAGAAAGAGAGAGAGAGA 1198 20 CAGAGAGAGAGAGAGA 1 Similarity Conservative 90.0%; 0, Score 16.8; Pred. No. 5. Mismatches 6e+02 DB 1; Length 21; Indels 0 Gaps 0

10

T; 0 U; 0 Other;

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Human hair keratin-associated-protein PCR primer P7
                                       06-MAY-2004
                                                                           ADJ92226;
                                                                                                              ADJ92226 standard; DNA; 21
                                     (first entry)
                                                                                                              ₽P
   SEQ ID NO:85
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22-MAY-2003 hair; keratin-associated protein; KAP; human; keratin; toiletry; therapeutic; hair growth promoter; hair disorder; PCR; primer; s WO2003042387-A1. Homo sapiens.

RESULT 425
ADJ92226
ID ADJ922
XX ADJ922
XX ADJ922
XX O6-MAY
XX Hair;
KW hair;
KW therap
XX Chomo s
OS Homo s
OS Synthe
XX Synthe
XX I ADJ922
XX I ADJ9 Kudo J, Shibuya K, 13-NOV-2001; 2001JP-00348050 (UYKB-) UNIV KBIO. (NIPR-) JAPAN SOC PROMOTION SCI 13-NOV-2002; 2002WO-JP011851 Shimizu N;

WPI; 2003-493307/46

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RESULT 426
ABD19062/c
ID ABD190
XX ABD190
XX BD190
XX Human;
XW Human;
XW Human;
XW respir
XW respir
XW surfac
XW semphys
XW emphys
XW emphys
XW gulmon
XX
BU HOMO &
XX
PM W02002
XX
PM W02002
XX
PM 31-OCT
XX
PF 23-APF
XX
PR 24-APF
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC the human and mouse DNA, and their partial peptides, and proteins derived carid residues; (3) DNA hybridising to the DNA encoding KAP; (4) peptides CC SCCXPSCCXP (I), where X = Q, V, R or I, and YGXXYYSGY (II), where X = Q, V, R or I, and YGXXYYSGY (III), where X = Y, CL or F; (5) fusion proteins and peptides containing these proteins and peptides together with a marker protein or peptide; (6) antibodies to the CC proteins and peptides; (7) recombinant proteins and peptides binding to the CC these antibodies; (8) expression vectors containing the DNA encoding KAP; (10) host cells transformed by these vectors; (10) non-human animals which CC are knockout animals for KAP or which overexpress KAP; (11) screening CC substances promoting or inhibiting the binding of KAP to hair keratin, or CC compositions on thair the expression of KAP; comprising using KAP or their partial peptides or cells expression them; (12) compounds CC compositions containing these compounds, or containing KAP or their CC compositions containing these compounds, or containing KAP or their CC compositions (such as hair and beard growth improvers, hair colourants and hair conditioners) and in therapeutic compositions for them to the present sequence is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
                                                                                                                                                                                                                                                  Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes DNAs encoding hair keratin-associated proteins (KAP) of human origin, which bind to hair keratin. Also described: (1) DNA encoding KAPs of mouse origin; (2) proteins encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNAs encoding 39 Keratin-associated proteins localized on human chromosome 21, useful for screening binding and expression modias cosmetic and therapeutic agents for hair disorders.
                       24-APR-2001; 2001US-0286036P
                                                                23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                                                                                                                                                                    Human histidine decarboxylase DNA fragment 1025.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABD19062;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABD19062 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; SEQ ID NO 85; 352pp; Japanese
                                                                                                                                                                                                                                 pulmonary transplantation rejection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3935 AGATCAACCCACAGCACAGC 3954
                                                                                                                                                                                        sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 BP; 7 A; 10 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGCTCAACCCCCAGCACAGC 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  股
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ŏ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       .6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (2) proteins encoded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         modifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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RESULT 427
AAQ25483
ID AAQ2548
XX AAQ254
AC AAQ254
XY 25-MAR
DT 25-MAR
DT 07-DEC
XX U7-DEC
XX Target
XW malign
XX malign
XX MO920:

rich

HUMINFAA target duplex sequence.

Target; human Tumour necrosis factor mRNA; AIDS; triplex; HIV; hepatitis;

25-MAR-2003 07-DEC-1992

(revised)
(first entry)

AAQ25483 standard;

DNA; 22

AAQ25483;

Ś

1179 CAGAGAAAGAGAGAGAGAGA 1198

Conservative

<u>,</u>

Mismatches

0

Gaps

0

90.0%; 0.3%;

Score 16.8; DB 1; Length 21; Pred. No. 5.6e+02;

Query Match
Best Local Similarity
Matches 18; Conserv

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CC of the invention has antiallergic, antiinflammatory, antiasthmatic, and the invention has antiallergic, antiinflammatory, antiasthmatic, activity, is a CC beta-adrenergic agonist. The composition is useful for preventing or CC treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production CC or availability, or to increase the degradation of the target mRNA or to CC reduce the amount of target polypeptide present in the lungs. The CC pulmonary obstruction, and/or bronchoconstriction and/or lung cinflammation, allergies and/or surfactant hypoproduction are associated CC with a disease or condition such as pulmonary vasoconstriction, conflammation, allergies, asthma, impeded respiration, respiratory CC inflammation, emphysema, chronic obstructive pulmonary disease, pulmonary CC transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to CC thymidines present in the target RNA serves to prevent the breakdown of CC thymidines present in the target RNA serves to prevent the breakdown of CC eng., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targe nucleic acids associated with lung airway or lung dysfunction,
Sequence 21 BP; 0 A; 10 C; 1 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            bronchoconstriction, respiratory tract inflammation, allerg reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comprising oligonucleotides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes a novel composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Li Y, Sa
Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 10404; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sandrasagra A, K
L, Shahabuddin S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              effective
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Katz
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H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for alleviating
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (a) a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       first active agent,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          allergies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   targeted to
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RESULT 428
AAT76387/c
ID AAT763
XX AAT763
XX IS-SEP
XX Human
XX Asthma
KW Chroni
XX Synthe
PN WO964(
XX PP 06-JUI
PR 07-JUI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    맑
                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17-APR-1991;
17-APR-1991;
27-SEP-1991;
                                                                                                                                                          Asthma;
chronic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequence depicts a HUMTNFAA sequence beginning at nucleeotide 1137. The sequence is a viral duplex sequence which contains a purine-rich region concentrated on one chain of the duplex. The sequence may be prepd. by standard DNA synthesis. The HUMTNFAA duplex sequence is used as a target for novel oligomers which are capable of forming a triplex at physiological ph by coupling into the major groove of the DNA duplex. Ten such oligomers TNF 211-20 are capable of forming a triplex with this sequence. The oligomers are used in the treatment of inflammation. Similar oligomers may be used to target viral DNA duplexes specific for HIV, herpes and other viruses. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. The oligomer is able to inhibit gene expression, as verified by in vitro systems. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                    Human tumour necrosis factor alpha antisense oligonucleotide HSTNFAAS6
                                                                                                                                                                                                                                       15-SEP-1997
    07-JUN-1995;
                                 06-JUN-1996;
                                                                19-DEC-1996.
                                                                                              WO9640162-A1
                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                   AAT76387;
                                                                                                                                                                                                                                                                                                 AAT76387 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 11 A; 0 C; 11 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 11; Page 64; 77pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New oligomers contg. modified bases - doublet in a DNA duplex, for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-APR-1991;
17-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-NOV-1990;
18-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               herpes malignancy and inflammation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1992-217083/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Froehler B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-JUN-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GILE-) GILEAD SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                          1181 GAGAAAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                             _
                                                                                                                                                        airway epithelium; adenosine free; cystic fibrosis; obstructive pulmonary disease; bronchitis; ss.
                                                                                                                                                                                                                                                                                                                                                                             GGGGAAGAGAGAGAGAAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Krawczyk S,
                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
91US-00766733.
    95US-00474497
                                   96WO-US009306
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90US-00617907
91US-00643382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91WO-US008811.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matteucci MD,
                                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16.8; D
Pred. No. 5.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 which form a triplex with G-C and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                    .7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Milligan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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ARSSULT 429
AAXS4536/C
ID AAXS45
XX AAXS45
XX ANTise
CKW impair
KW impair
KW pulmon
KW respir
KW pulmon
KW coloroni
KW prosta
XX prosta

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A method for treating airway disease in a subject has been produced, which involves the topical administration of an essentially adenosine free antisense oligonucleotide (ON) to the airway epithelium of the subject. The present sequence is an antisense oligonucleotide HSTWFAAS6 specific for the human tumour necrosis factor alpha. The method can be used to treat airway diseases such as cystic fibrosis, asthma, chronic obstructive pulmonary diseases bronchitis and other airway diseases characterised by an inflammatory response. By eliminating adenosine from the antisense ON, its liberation upon antisense degradation is prevented, thereby preventing adenosine- induced bronchoconstriction in patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      acute asthma; allergy; asthma; impeded respiration; respiratory distress syndrome; pain; cystic fibrosis; pulmonary hypertension; pulmonary vasconstriction; emphysema; chronic obstructive pulmonary disease; leukemia; lymphoma; carcinoma; colon cancer; breast cancer; lung cancer; pancreatic cancer; hepatic metastasis;
                                                                                                                                                                                    17-SEP-1997;
09-JUN-1998;
                                                                                                                                                                                                                                                                                                                                           25-MAR-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            prostate cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense oligonucleotide; multiple target; antisense treatment; impaired respiration; inflammation; lung disease; pulmonary vasoconstriction; inflammation; allergic rhinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human adenosine Al receptor antisense oligonucleotide fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-JUL-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX54536 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 0 A; 11 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           thereby preventing adenosine-
with hyper-reactive airways
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 5; Page 37; 71pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Treatment of airway diseases such as asthma - by topically applying adenosine-free antisense oligo:nucleotide to airway epithelium of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-051871/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Мусе ЈЖ,
                                                                                                                                                                                                                                                                              17-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                     WO9913886-A1
                                                                                                                       (UYEC-) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (UYEC-) UNIV EAST CAROLINA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1181 GAGAAAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22 GGGGAAGAGAGAGAGAAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       h 0.3%;
Similarity 90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                          BAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                 97US-0059160P
98US-00093972
                                                                                                                                                                                                                                                                              98WO-US019419
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16.8;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; DB 1;
5.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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WPI; 1999-229400/19

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RESULT 430
AAA33980/c
ID AAA339
XX AAA339
XX AAA339
XX Low ad
XX Human;
KW Phosph
KW Phosph
KW allerg
KW lung allerg
KW lung allerg
KW lung blmon
KW cancer
XX Homo s
PN WO2000
XX Homo s
PN WO2000
XX O3-AUG
PF 03-AUG
XX O3-AUG
PF 03-AUG
XX O3-AUG

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 18; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            diseases and conditions. Typical diseases and conditions are those associated with impaired respiration and inflammation, including lung diseases, pulmonary vasoconstriction, inflammation, altergic rhinitis, acute asthma, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, pulmonary hypertension, pulmonary vasoconstriction, emphysema, chronic obstructive pulmonary disease (CODD), and cancers such as leukemias, lymphomas, carcinomas e.colon cancer, breast cancer, lung cancer, pancreatic cancer, hepatocellular carcinoma, kidney cancer, melanoma, hepatic metastases, a well as all types of cancers which may metastasize or have metastasized
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   directed against at least 2 mRNAs selected from target genes, coding and non-coding regions of RNAs corresponding to target genes, gene initiation codons, genomic flanking regions, intron-exon borders, the 5'-end, the 3'-end and the juxta-section between coding and non-coding regions and all segments of RNAs encoding proteins associated with one or more diseases, conditions or mixtures. The antisense oligonucleotides may be derived from sequences AAX55272-74. These multiple target oligonucleotides (specifically AAX55180-271) can be used for the antisense treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              phosphorothicate; impaired respiration; inflammation; allergy; allergy; disease; bronchoconstriction; inhibitor; antiinflammatory; antiallergic; antiasthmatic; cytoostation analgesic; impaired airway; lung disease; ischaemic condition; pulmonary vasoconstriction; asthma; respiratory distress syndrome; pain; cystic fibrosis; emphysema; pulmonary hypertension; chronic obstructive pulmonary disease; COPD;
                                  WPI; 2000-205971/18
                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200009525-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; adenosine receptor; low adenosine antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAA33980 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22 BP; 0 A; 11 C; 0 G; 11 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides used in treatment of, e.g. pulmonary vasoconstriction.
                                                                                                                                                                                                                                                                                                        03-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Low adenosine antisense oligonucleotide SEQ ID NO:1669.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       to the lungs,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 27; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The specification describes antisense oligonucleotides (AAX52869-X55271)
                                                                                                                                                                        (UYEC-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1181 GAGAAAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22
                                                                                                                                                                        VIND
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGGGAAGAGAGAGAGAAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                        EAST CAROLINA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       including breast and prostate cancer
                                                                                                                                                                                                                                          98US-0095212P
                                                                                                                                                                                                                                                                                                        99WO-US017712.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      lymphoma; carcinoma; metastasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16.8;
No. 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28
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The present invention describes a new composition comprising an antisense CC oligonucleotide (ON) with low adenosine (up to 15%), which targets cnucleic acids involved in bronchoconstriction, allergies, and/or CC inflammation. The ON can have antiinflammatory, antiallergic, antiasthmatic, cytostatic and analgesic activities. The compositions are CC useful for the treatment of diseases associated with inflammation, CC impaired airways, including lung disease and diseases whose secondary CC effects afflict the lungs of a subject. They can be used for treating CC e.g. ischaemic conditions, pulmonary vasoconstriction, allergies, asthma, cimpaired airways, pulmonary vasoconstriction, allergies, asthma, CC impaded respiration, respiratory distress syndrome, pain, cystic CC fibrosis, pulmonary hypertension, emphysema, chronic obstructive CC pulmonary disease (COPD), and cancers such as leukaemias, lymphomas, CC carcinomas, and cancers which may metastasise to the lungs, including breast and prostate cancer. The reduction of the adenosine content of the CC NS reduces side effects. The A-containing ONs break down with the CC uncleotide sequences given in the sequence listing from the present the mucleotide sequences given in the sequence listing from the present the present invention, which correspond to SEQ ID NO:1 to 2815, and then the last 185 cquences given in the disclosure of the present invention do not match CC up with their corresponding SEQ ID NO: sequences given in the sequences given in th
Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotides useful for treating e.g. pulmonary vasoconstruction, inflammation, allergies, asthma, hypertension, bronchitis, emphysema, respiratory distress syndrome, ischemia or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 18; Page 472; 1343pp; English.
   BP; 0 A;
   11 C; 0
   G; 11 T;
       0 U;
       0 Other;
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ARESULT 431
AAF20102/C
ID AAF2010
XX AAF201
XX AAF201
XX AAF201
XX Low ac
KW Low ac
KW human;
KW surfac
KW respin
KW respin
KW respin
KW respin
KW respin
KW chromo;
KW chromo;
KW chromo;
KW chromo;
XX CACCE
XX CACCE
XX CACCE
XX WO2000
XX PD 26-OC;
XX Y
                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                                                                                                                                                                                                                                 AAF20102;
                                                                                                                                                                                                                                                   AAF20102 standard; DNA;
                                                                                                                                                                                                               14-MAR-2001
                                                                                                                                                                                                                                                                                                                     1181 GAGAAAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                   GGGGAAGAGAGAGAGAAA 3
                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                0.3%;
                                                                                                                                                                                                                                                   22
                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                Score 16.8;
Pred. No. 5.
                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                              .7e+02;
                                                                                                                                                                                                                                                                                                                                                         1; Length 22;
                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                                                                                       Gaps
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human; airway disorder; bronchoconstriction; lung inflammation; surfactant depletion; respiratory; bronchodilator; antiinflammatory; immunosuppressive; antiasthmatic; analgesic; hypotensive; cytostatic; respiratory obstruction; pulmonary obstruction; impeded respiration; surfactant hypoproduction; pulmonary vasoconstriction; asthma; RDS; respiratory distress syndrome; pain; cystic fibrosis; allergic rhinitis; pulmonary hypertension; emphysema; pulmonary transplantation rejection; pulmonary hypertension; emphysema; pulmonary transplantation rejection;

24-MAR-2000; 2000WO-US008020.

26-OCT-2000 WO200062736-A2

88.

obstructive

pulmonary disease;

pulmonary infection; bronchitis,

Human tumour necrosis factor alpha polynucleotide fragment #1669

Low adenosine antisense oligonucleotide; phosphorothicate; allergy;

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RESULT 432
ACC85489/c
ID ACC854
XX ACC854
XX ACC854
XX ACC854
XX LACC854
XX Late F
XX ACC854
XX MO2003
XX MO2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                  S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC immunosuppressive, antiasthmatic, hypotensive and cytostatic activities.

CC The antisense oligonucleotides and (I) can be used to down-regulate the cc expression and or activity of target polypeptides associated with cc lung/respiratory disorders and malignancies, such as stimulating and activating peptide factors and transmitters, transcription factors, crimmunoglobulins and antibodies, antibody receptors, cytokines and cc enhanckines, endogenously produced specific and non-specific enzymes, binding proteins, adhesion molecules and their receptors, cytokine and cc chemokine receptors, adenosine receptors, bradykinin receptors, cytokine and craceptors, cytokine and craceptors, cytokine and craceptors, cytokine and craceptors, binding proteins and malignancy associated proteins. The antisense oligonucleotides may be used in this way to treat disorders including respiratory obstruction (especially pulmonary obstruction and/or lung inflammation, allergy(ies) and/or sonchance condition selected from pulmonary vasoconstriction, inflammation, condition selected from pulmonary vasoconstriction, inflammation, callergies, asthma, impeded respiration, respiratory disease or condition selected from pulmonary vasoconstriction, inflammation, condition, camplysema, chronic obstructive pulmonary disease (COPD), pain, cystic fibrosis (CF), allergic rhinitis (AR), pulmonary cancer. AAF18434 to AAF21543 represent human polynucleotide from the exemplification of the present human polynucleotide condition of the present human polynucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches 18;
                                                                                                  Late PCR; polymerase chain reaction; human; CFTR; HEX-A; limiting primer; excess primer; PCR; primer; ss.
                                                                                                                                                                                                                                                                                           ACC85488 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22 BP; 0 A; 11 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes low adenosine (A) content antisense oligonucleotides and compositions (I) comprising them. In the antisense oligonucleotides the A is replaced by a 'Universal' or alternative base. (I) can have respiratory, bronchodilator, antiinflammatory, analgesic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 14; Page 241; 1592pp; English.
                       WO2003054233-A1
                                                              Homo sapiens
                                                                                                                                                              Human HEX-A gene PCR primer SEQ ID NO:
                                                                                                                                                                                                         29-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        adenosine receptors during metabolism, useful e.g. for treating cancers and respiratory obstructions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-679539/66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Low adenosine (A) content antisense oligonucleotides which do not trigger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NYCE/) UNIV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                        1181 GAGAAAGAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                                                GGGGAAGAGAGAGAAA 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EAST CAROLINA.
J W.
                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99US-0127958P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8; DB 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Best Local (
              Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a method of non-symmetric polymerase chain reaction (PCR) amplification, comprising thermally cycling a PCR reaction mixture containing a DNA amplification target sequence, a pair of PCR primers, dNTP's and thermostable polymerase repeatedly through PCR steps of strand melting, primer annealing, and primer extension. The method is useful for amplifying stretches of DNA, including cDNA reverse transcribed from RNA, for assays, for diagnostics and other purposes. The present sequence is a primer used to demonstrate the method of the
                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Non-symmetric polymerase chain reaction amplification, useful for amplifying DNA by thermally cycling a PCR reaction mixture containing DNA amplification target sequence, a pair of PCR primers, dNTP's and
                                                                                     24-APR-2001; 2001US-0286137P
                                                                                                               23-APR-2002; 2002WO-US013135
                                                                                                                                                                                                                                             antisense gene therapy; respiratory; lung; adenosine sensitivity;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                  Human tumour necrosis factor antisense fragment no.1660
                                                                                                                                                                                                                                                                                                                                                              17-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                     ABZ95796 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 22 BP; 0 A; 11 C; 4 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 66; 125pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             thermostable polymerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-DEC-2001; 2001US-0341886P.
17-DEC-2002; 2002US-00320893.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-DEC-2002; 2002WO-US040752.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-JUL-2003
                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (UYBR-) UNIV BRANDEIS
                                                                                                                                                                                                                               inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        909 CCAGGGCTCAGAGAGAAGGG 928
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18;
                                                          EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CCAGGGGGCAGAGAGAAGGG
              Li Y,
Tang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pierce K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                              (first entry)
              Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                       22
                                                                                                                                                                                                                                                                                                                                                                                                                       BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 16.8; DB 1;
Pred. No. 5.7e+02;
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                             Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rice J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sanchez JA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 22;
                               Aguilar
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RESULT 434
ABD19536/c
ID ABD195
XX ABD195
XX ABD195
XX Human
DE Human
XX Human;
KW respix
KW respix
KW analge
KW analge
KW pulmor
XX Demons
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC has antiinflammatory, antiallergic, antiasthmatic, hypotensive, CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or CC preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapeutic respiratory effect of an CC antiinflammatory steroid in a subject, for reducing or depleting levels CC for reducing sensitivity to adenosine, reducing levels of adenosine crecaptor, producing bronchodilation, increasing levels of adenosine crecaptor, producing bronchodilation, increasing levels of biquinone or CC lung inflammation, lung allergies, or a respiratory disease or condition. CC Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO CC at fip.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 18; Conser
                                                                                                                                                                                                                                                                                                      Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human tumour necrosis factor DNA fragment 1660.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                        24-APR-2001; 2001US-0286036P
                                                                                                                                   23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                       WO200285309-A2
                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ABD19536;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABD19536 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonucleotide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 11038; 872pp; English.
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                                             (EPIG-) EPIGENESIS PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
Sandrasagra A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
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90.0%;
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Pred. No. 5
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  Katz
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Aguilar
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expression of a target polypeptide associated with lung airway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA.

C The invention also describes a kit, that comprises: (a) a delivery containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, canalgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or composition comprises oligo and is administered to reduce the amount of target polypeptide present in the lungs. The pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascomstriction.

C inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascomstriction.

C inflammation, allergies, asthma, impeded respiration, respiratory chapteress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system c.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
  Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptor surfactant depletion or hyposecretion, when administered to a mammal oligonucleotides are derived from a gene encoding or regulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bronchodilating agent.
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BP; 0 A; 11 C; 0 G; 11 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    763pp; English
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a mammal. The
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RESULT 435
AAA29753/c
ID AAA297
XX AAA297
XX AAA297
XX 15-AUG
XX 15-AUG
DE Synthe
XX Primer
XW Primer
XW purifi
XX Synthe
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Best Local Similarity
Matches 18; Conserv
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                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                     purification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic oligonucleotide #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-AUG-2000
                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                        Primer; destabilise non-specific duplex formation; PCR; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA29753 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                     sequencing; genetic marker; RACE;
                                                                                                                                     Location/Qualifiers
/mod_base= i
/note= "inosi
                                                                                       /*tag= a
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                                                                                                                                                                                                                                                                                                                                                     DNA synthesis;
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ARESULT 436
AAC83384
ID AAC833
XX AC833
XX Primer
XX Primer
XX Primer
XX Uniden
XX W02000
XX 09-NOV
XX 28-APR
XX 28-APR
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a method for destabilising non-specific duplex formation, between an oligonucleotide and a target nucleic acid (NA), comprising incubating the target NA with a modified oligonucleotide (I) comprising incubating the target NA with a modification which decreases or abrogates H-bonding between the modified oligonucleotide and the non-specific target NA. The modified oligonucleotide is used to improve discrimination between the targeted homopolymeric sequence and a non-homopolymeric target sequence. It is used to increase the proportion of full length cDNA clones for a library, to reduce misprining during sequencing, 5, or 3, RACE (rapid amplification of CDNA ends) or DNA synthesis or to generate bona fide genetic markers. The present sequence represents an oligonucleotide which is used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Novel method of stabilizing duplex formation, or destabilizing non-
specific duplex formation using primer containing modified nucleotide
analogs, useful for preventing mispriming during PCR, RACE, DNA synthesis
                                                                                                                       Unidentified
                                                                                                                                                 Primer;
                                                                                                                                                                            Primer -CS
                                                                                                                                                                                                       26-FEB-2001
                                                                                                                                                                                                                                 AAC83384;
                                                                                                                                                                                                                                                            AAC83384 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 23 BP; 0 A; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-OCT-1998;
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               28-APR-1999;
                                        28-APR-2000; 2000WO-GB001639
                                                                                              WO200066774-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequencing.
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                                                                                                                                                                                                                                                                                                                               23
                                                                                                                                                                                                                                                                                                                                                                                   18;
                                                                                                                                                  virus;
                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                             (outer).
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                                                                                                                                                  resistance;
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81.8%;
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                                                                                                                                                                                                                                                            24 BP
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Pred. No. 5.
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                                                                                                                                                                                                                                                                                                                                                                                             .7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
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present invention

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Best Local Similarity
Matches 18; Conserv
                                                                               The present invention describes human carbamylaspartic dehydrase 9.46 (I). Also described is a DNA recombination process used to produce (I) (I) can be used in the treatment of various diseases, such as malignan tumours, haemopathy, HIV infection, immunological diseases and various inflammations. The present sequence represents a PCR primer for (I),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to detecting viruses resistant to antiviral drugs. The invention involves recombination of a DNA vector encoding a modified virus with nucleotide sequences from virus samples
                                                                                                                                                                                                                                                  Example 2; Page 16 (Disclosure); 33pp; Chinese.
                                                                                                                                                                                                                                                                                                   New polypeptide-human carbamylaspartic dehydrase encoding the polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Mao Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-NOV-2000; 2000CN-00127141.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-NOV-2000; 2000CN-00127141.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-JUN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; carbamylaspartic dehydrase 9.46; enzyme; malignant tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human carbamylaspartic dehydrase 9.46 PCR primer 2 SEQ ID NO:4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABQ82590;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             suspected of being drug-resistant, to produce recombinant viruses
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 13; 36pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Assay for detecting viruses resistant to anti-viral drugs by recombination of DNA vector comprising a wild type virus strain genome
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        naemopathy; HIV infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              immunological disease; inflammation;
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Pred. No. 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SHANGHAI.
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5.7e+02;
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                                                                                                                                                                                                                                                                                                                                  polynucleotide
                                                                                                                                           as malignant
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RESULT 439
ACC85493/c
ID ACC854
XX
AC ACC854
XX
DT 29-SEP
XX
DE Human
XX
KW Late F
KW excess
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ABQ79142/c
ID ABQ791
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Best Local (
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Best Local
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 Late PCR; polymerase chain reaction; human; excess primer; PCR; primer; ss.
                                                                                                                                                                                                                                                              Sequence 24 BP; 4 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                   This invention relates to a novel polypeptide-human eukaryotic initial factor -2 (elf-2)-associated protein 74.14 and the application of the polypeptide in treating diseases such as tumours, virus infection abnormalities and development disorders. The present sequence represents a primer related to the novel human eukaryotic initial factor -2 (elf-2)-associated protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; eukaryotic initial factor -2; (elF-2); tumour; virus infection; development disorder; PCR; primer; ss.
                                   Human CFTR gene PCR primer SEQ ID NO:
                                                           29-SEP-2003
                                                                                   ACC85493
                                                                                                          ACC85493 standard;
                                                                                                                                                                                                                                                                                                                                                                     Example
                                                                                                                                                                                                                                                                                                                                                                                        Polypeptide-human eukaryotic initiation factor-2 (eIF-2)-associated protein 74.14 and polynucleotide encoding it.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-MAR-2002.
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21 ATTAMAMATTTTTTATATAA 2
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                                                           (first
                                                                                                        DNA;
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                                                                                                                                                                                                                         Score 16.8; DB 1;
Pred. No. 5.7e+02;
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Pred. No. 5
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                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
            CFTR; HEX-A; limiting primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2
                                                                                                                                                                                                                                      Length 24;
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RESULT 440
ADG35077/c
ID ADG350
XX ADG350
AC ADG350
XX Human
XX Hu
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Best Local S
Matches 18
                                                                                                                                                                                                                                                        RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; tumour necrosis factor receptor; TNF receptor; human; DNA-RNA hybrid; antibacterial; immunosuppressive; antirheumatic; antiarthritic; antiantipsoriatic; antiinflammatory; septic shock; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to a method of non-symmetric polymerase chain reaction (PCR) amplification, comprising thermally cycling a PCR reaction mixture containing a DNA amplification target sequence, a pair of PCR primers, dNTP's and thermostable polymerase repeatedly through PCR steps of strand melting, primer annealing, and primer extension. The method is useful for amplifying stretches of DNA, including cDNA reverse transcribed from RNA, for assays, for diagnostics and other purposes. The present sequence is a primer used to demonstrate the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Non-symmetric polymerase chain reaction amplification, useful for amplifying DNA by thermally cycling a PCR reaction mixture containing DNA amplification target sequence, a pair of PCR primers, dNTP's and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human TNF receptor siNA oligonucleotide SEQ ID NO:429.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-DEC-2001; 2001US-0341886P.
17-DEC-2002; 2002US-00320893.
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                                                                                                                                                                                                                                   HIV/AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 4; Page 72; 125pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
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                                                                                                                                                                                                                                      psoriasis; inflammation; autoimmune disease; target sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 0 A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90.0%;
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Pred. No. 5.7e+0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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. 5.7e+02;
2;
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                                                                                                                                                                                                                                                                                                                       DNA-RNA hybrid;
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WO2003070897-A2

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RESULT 441
ADK97129/c
ID ADK97129 standard; DNA; 23 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to short interfering nucleic acids (siNA) which CC downregulate expression of the human tumour necrosis factor (TNF) CC receptor gene by RNA interference. The siNAs may or may not comprise CC ribonucleotides and may be double or single stranded. They further CC comprise enese and antisense regions, or alternatively are assembled from CC a sense oligonucleotide and an antisense oligonucleotide. Specifically, CC micro-RNA (miRNA) and short hairpin RNA (shRNA), double-stranded RNA, CC micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be CC unmodified or chemically synthesised, expressed from a vector or enzymatically can be chemically synthesised, expressed from a vector or enzymatically convivo delivery of siNA, conjugates and/or complexes of siNA; and vectors that express siMA. The siNAs are used to modulate expression of the TNF complexes of siNA; and vectors converted of the expression of the TNF complexes of siNA; and vectors converted of the expression of the TNF complexes of siNA; and vectors converted of the expression of the transplants for the treatment of a converted of the expression of the transplants for the treatment of a converted of the expression of the treatment of a converted of the expression of the 
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Matches 19; Conserva
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11-MAR-2002; 2002US-036314P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
28-NOV-2002; 2002US-0429359P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 23 BP; 4 A; 6 C; 4 G; 0 T; 9 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New short interfering nucleic acid, useful e.g. for diagnosis of septic shock or rheumatoid arthritis, expression of the tumor necrosis factor gene.
                                                             human; single nucleotide polymorphism; SNP; ss; primer
                                                                                                         Primer of
                                                                                                                                                    06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-697609/66.
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                                                                                                                                                                                                                                                                                                                                                                                        4317 CTGCTATCGAAGCCCTGGAGAAG 4339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                    (first entry)
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                                                                                                         invention #2849.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     82.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 16.6; DB 1;
Pred. No. 6.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                  4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           for treatment and
is, downregulates
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                  <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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RESULT 442
AAV21969/c
ID AAV219
XX AAV219
XX Nuclea
XX Nuclea
XX Nuclea
XX Veteri
KW Veteri
KW Veteri
KW WO9803
XX WO9803
XX GOLIG-
XX (OLIG-
XX Arrow
XX WIF 23-JUL
XX Arrow
XX IT Arrow
XX IT Arrow
XX IT Arrow
XX IT Arrow
XX IT 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a polynucleotide isolated from a human gene and is useful for detecting a single nucleotide polymorphism in a human gene or for diagnosing of disease. The invention enables the detection of a single nucleotide polymorphism in a human gene. The present sequence represents a primer of the invention.
                                                                                                                 Treating bacterial infections in humans or animals with oligo:nucleotide(s) - resistant to nuclease and targette nucleic acid or proteins, also conjugates of these oligo with antibiotics.
                                                                                                                                                                                                                                                              WPI; 1998-120687/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                         24-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             veterinary medicine; treatment; human; industrial bacterial control; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nuclease resistant; bacterial infection; antibiotic; target;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nuclease resistant antisense oligo NBT 142 targeted against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV21969 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 2; SEQ ID NO 6158; 2627pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel polynucleotide useful for PCR amplification along with two DNA fragment from another set of sequences, or for detecting single nucleotide polymorphism in human gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-MAR-2002; 2002JP-00064373
                                                                                                                                                                                                                                                                                                                                                                                (OLIG-) OLIGOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2662 CACTCTCTGGAGTCCCTCCACTG 2684
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23 BP; 7 A; 4 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                    Dale RMK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                ETC & OLIGOS THERAPEUTICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                         96US-00685575
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97WO-US012961.
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                                                                                                                                                                                                                                                                                                                 Thompson
                                                                                                                                         resistant to nuclease and targetted to bacterial ins, also conjugates of these oligo:nucleotide(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 16.6; DB 1
Pred. No. 6.1e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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This antisense oligonucleotide is nuclease resistant and

can be used in

Claim 49; Page 87; 163pp; English

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RESULT 443
AAX91065/c
ID AAX910
XX AAX910
XX AAX910
XX AAX910
XX Phosph
KW Phosph
KW Dindir
XX Synthe
XX US5955
XX US5955
XX 16-NON
PR 21-SEE
XX 16-NON
PR 21-NON
XX (GENT-
XX Schwal
XX Schwal
XX WFI;
CC Inter
CC inter
CC inter
CC pure
CC pure
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   method is used to treat infections by a wide variety of Gram-positive and Gram-negative, or acid-fast, bacteria, in human and veterinary medicine. The methods are particularly used in immuno-compromised individuals (e.g. patients with acquired immunodeficiency syndrome or those receiving chemotherapy or radiation therapy; optionally in combination with, or fused to, antiviral or other antimicrobial oligonuclectides. Apart from therapeutic use, the oligonucleotides can be used to control bacteria in laboratory cultures, foods, beverages and industrial processes. The oligonucleotides are specific for bacteria, without affecting metabolism in mammalian cells. They may also activate Russe H and have a general,
The invention provides methods for preparing oligomers having phosphonate internucleosidyl linkages of a preselected chirality which hybridize to a target RNA sequence. The method of making comprises: (a) synthesizing a nucleoside dimer, trimer, or tetramer with racemic internucleosidyl phosphonate linkages; (b) purifying the racemic nucleoside to a chirally pure nucleoside; and (c) sequentially linking at least 2 of the chirally pure nucleosides to form a synthetic oligomer that is enriched for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             non-specific immune-stimulating effect. The oligonucleotides can be administered orally, intranasally, rectally, topically or by injection, optionally coupled to an agent (e.g. carbohydrate or polyamine) that
                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           and formulated with a carrier. A compound comprising this nuclease resistant oligonucleotide can be covalently linked to an antibiotic. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and formulated with a carrier.
                                                                                                                                                                                                                          Oligomers
                                                                                                                                                                                                                                                                  WPI; 1999-539600/45.
                                                                                                                                                                                                                                                                                                                                                                                        21-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                         16-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        binding affinity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Phosphonate internucleosidyl linkage; chirality;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CAT gene target RNA fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-NOV-1999
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                                                                                                                                                                                                                                                                                                                                                 GENT-) GENTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
mes 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 treatment of animals, including humans, having a bacterial infection treatment comprises administration of such nuclease resistant gonucleotides, targeted to a nucleic acid or protein of the bacterium,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1181
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                                                                                                                                                                                                                                                                                                         DA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cellular uptake
                                                                                                                                                                                                       made
with
                                                                                                                                                                Col 41-42; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                 INC.
                                                                                                                                                                                                                                                                                                     Vaghefi
                                                                                                                                                                                                   using chirally pure nucleoside dimers, trimers, enhanced binding affinities.
                                                                                                                                                                                                                                                                                                                                                                                      93US-00154013
94US-00343018
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.3%;
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                                                                                                                                                                                                                                                                                                       Riley
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                                                                                                                                                                                                                                                                                                         TA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16.4; DB 1,
No. 6.3e+02;
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                                                                                                                                                                                                                                                                                                       Arnold LJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hybridization;
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                                                                                                                                                                                                                                                                                                         MA;
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RESULT 444
ADH70321
ID ADH70321
AC ADH703
AC A
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Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; Addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; degenerative nervous system disease; multiple sclerosis; Alzheimer's disease; hypersensitivity disease; type I hypersensitivity; allergy; type II hypersensitivity; leprosy; infectious disease; viral infection; type IV hypersensitivity; leprosy; infectious disease; viral infection; HIV; fungal infection; Candida; parasitic infection; schistosome; filaria; bacterial infection; Mycobacterium; neoplastic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18
                                                                                                                                                                                                                                                               Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, nucleic acid primers specifically priming and allowing amplific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human Vbeta gene repeat sequence #111.
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                                                                                                                                                                                                                                                                                                                                                                                                                           Hood LE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-SEP-1994;
19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (ROWE/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (\dooh/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ymphoproliferative disease; leukaemia; lymphoma; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
                                                                                                                                                                                                                                         gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HOOD L E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ROWEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AGAGAAAGAGAGAGAG 1197
                                                                                                                                                                                                                                                                                                                                                                                                                           Rowen L;
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95US-00531241.
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0; Mismatches
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Pred. No. 6
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                                                                                                                                                                                                                                                                    disease, compa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     brain cancer
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The invention relates to a kit for diagnosing and treating associated diseases which comprises a panel of nucleic acions.

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515; 164pp;

English

associated diseases which comprises a panel of nucleic acid primers specifically priming and allowing amplification of each Vbeta gene, VbetaRNA or CDNA. The kit is useful for diagnosing organ transplant rejection and diagnosing and treating T-cell associated diseases

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RESULT 445
ADH70679
ADH70679
ADH70679
ACC ADH706
XX ADH706
XX Human
XX Human
XX Human
XX Human
XX Addi:
XW Cype
XW HIV;
XW Eil:
XW Eil:
XW Eil:
XW Eil:
XW Eil:
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infections disease; neoplastic disease; hypersensitivity disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; degenerative nervous system disease; multiple sclerosis; hzpersensitivity disease; type I hypersensitivity disease; type I hypersensitivity; allorsy; type II hypersensitivity; leprosy; infectious disease; viral infection; type IV hypersensitivity; leprosy; infectious disease; viral infection; filaria; bacterial infection; Mycobacterium; neoplastic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sclerosis and Alzheimer's disease. Hypersensitivity diseases include Type I hypersensitivities such as contact with allergens that lead to allergies. Type II hypersensitivities such as those present in Goodpasture's syndrome and Type IV hypersensitivities such as those manifested in leprosy. Infectious diseases include viral infections caused by viruses such as HIV, fungal infections such as those caused by the yeast genus Candida, parasitic infections such as those caused by schistosomes, filaria and bacterial infections such as those caused by Mycobacterium. Neoplastic diseases include lymphoproliferative diseases such as leukaemias, lymphomas and cancers such as cancer of the brain, breast. The present sequence represents a Vbeta gene repeat sequence.
                                                                                       Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, compuncteic acid primers specifically priming and allowing amplification
                                                                                                                                                                                                                                                                                                                                                                                                                            19-SEP-1994;
19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human Vbeta gene repeat sequence #469.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 18
                                                                                                                                                                                                                 WPI; 2004-059052/06
                                                                                                                                                                                                                                                                         Hood LE, Rowen L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-OCT-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US2002150891-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADH70679 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        graft versus host disease, hypersensitivity diseases, infection and neoplastic diseases. Autoimmune diseases include Addison's
                                                                                                                                                                                                                                                                                                                                    (HOOD/) HOOD L E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .ymphoproliferative disease; leukaemia; lymphoma; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1180 AGAGAAAGAGAGAGAGAG 1197
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                cancer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGAGAGAGAGAGAGAG 18
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95US-00531241.
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Pred. No. 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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Disclosure; SEQ

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873; 164pp; English

primers AAV30470-V30490 were used for sequencing the coding region of the canine beta 3-adrenergic receptor (RA-Ca-b3) gene (AAV30469). RA-Ca-b3 has been implicated in obesity and obesity-related metabolic disorders e.g. diabetes. The canine version of RA-Ca-b3 can be used to develop

gene.

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RESULT 446
AAV30490
ID AAV304
XX AAV304
XX Canine
XX Canine
XX Canine
XX Canine
XX Canine
XX Canis
OS Canis
PN W09735
XX (VETI-
XX (VETI-
XX WPI;
PT Lenzer
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PT Useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                       Canine beta 2 and beta 3 adrenergic receptors useful for identifying specific ligands and (specific treatments for obesity in dogs.
                                                                                                                                                                                                                                                WPI; 1998-032136/03.
                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Canine; beta-adrenergic receptor; brown adipose tissue; hybridisation; ligand; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Canine beta-3 adrenergic receptor sense primer RT28.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAV30490 standard; DNA; 19
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                                                                                                                                                                                                                                                                                       Lenzen G,
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                                                                                                                                                                                                                                                                                                                                        (VETI-) VETIGEN
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17; Conserv
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                                                                                                                                                                                                                                                                                            Pietri-Rouxel
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                                                                                                        79pp;
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Pred.
                                                                                                                                                                                                                                                                                            Drumare
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No. 6.3e+02;
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                                                                                                                                                                                                                                                                                            Strosberg
                                                                                                                                                                              s and coding sequences -
(ant)agonists to develop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 18;
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RESULT 447
AAH45567/c
ID AAH455
XX AAH455
XX AAH455
XX Peroxi
KW Peroxi
KW Peroxi
KW PAPP
XX JP2001
XX JP2001
XX JP2001
XX JP201
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RESULT 448
ABA82228/c
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AC ABA822
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                                                                                                                                                                                                                                                             callus. Included in the invention are the gene encoding the peroxidase, vector containing the gene, and a method for the preparation of the peroxidase. The gene is useful for the development of a plant highly resistant against active oxygen which is formed under various environmental stress conditions. This sequence represents a PCR primer used in the isolation of peroxidase related DNA fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               treatments specific for dogs. differential screening for liq adrenergic receptor (AAW44932)
                ABA82228;
                                                                                                                                                                                                                                     Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JP2001095585-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Peroxidase; active oxygen resistance; transgenic environmental stress; ss.
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                                             ABA82228 standard;
                                                                                                                                                                                                                                                                                                                                                       This invention relates to a peroxidase derived from a Paraquat resistant
                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 20; 23pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                          Peroxidase derived from Paraquat-resistant callus, and gene encoding used for the development of plants resistant to active oxygen formed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-SEP-1999;
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                                                                                                                                                                                            Local
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                                                                                                                                                                                                                                    BP; 4 A; 7 C; 3 G; 5 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 5 A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                             Conservative
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                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                 stress.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              for dogs. The sequence can also be used in ing for ligands for RA-Ca-b3 as compared to the beta-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
                                                                                                                                                                                         94.4%;
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Pred. No.
                                                                                                                                                                                            Pred.
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No. 6.3e+02;
                                                                                                                                                                                            No.
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                                                                                                                                                                                                                                     0 Other;
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RESULT 449
ABK23025/c
ID ABK2300
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AC ABK230
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DT 09-APR
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DE Human
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Human
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Human;
KW Human;
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KW osteop
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Best Local Similarity
Matches 17; Conserv
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ABAB2038 to ABA82700 and AAG68168 to AAG68193 represent sequences used in the exemplification of the present invention
                                                                                       Human; mouse; Zmax1; HBM; high bone mass gene; lipid regulation; stroke; lipid-associated condition; arteriosclerosis; cardiovascular disease; ss; osteoporosis; atherosclerosis; diabetic atherosclerosis; plaque build-up; neurovascular condition; wound healing; gene therapy; PCR primer; probe; bone development disorder; antiarteriosclerotic; cardiovascular;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; high bone mass; HBM gene; Zmaxi gene; chromosome 11; 11q13.3; sequence tagged site; STS; osteoposis; osteopathic; gene therapy; antisense therapy; vaccine; bone disorder; Paget's disease; adapter; sclerostosis; osteomalacia; fibrous dysplasia; PCR primer; linker; ss
                                                                                                                                                                              Human Zmax1
                                                                                                                                                                                                             09-APR-2002
                                                                                                                                                                                                                                           ABK23025;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 34; 443pp; English.
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05-APR-2000;
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                                             Homo sapiens.
                                                                                                                                                                                                                                                                         ABK23025
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                                                                       osteopathic; cerebroprotective.
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                                                                                     development
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                                                                                                                                                                                                                                                                                                                                                                  GTTCCTGGGTGAGATCAA 3941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 bone mass (HBM) and ng bone mass for the
                                                                                                                                                                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                                                                GTTCCTGGGCGAGATCAA 2
                                                                                                                                                                                cDNA forward PCR primer #94.
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                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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2000US-00544398.
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Pred. No. 6.3e.
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                genes and proteins useful
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RESULT 450
ACC45608/c
ID ACC456
XX ACC456
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XX Human
XX Human
XX Gene t
KW Gene t
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XX I1-MA)
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Best Local
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                                                                                                                                                                                                                                                        Human; high bone mass; HBM; LRP5; LRP6; transgenic; bone mass modulation; gene therapy; bone density modulation; bone etrength; trabecular number; bone size; bone tissue connectivity; bone disease; osteoporosis; PCR; osteomalacia; rickets; Paget's disease; neoplasm of the bone; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; Page 39; 409pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying molecules involved in lipid regulation, useful for diagnosing, treating or preventing e.g., arteriosclerosis, comprises identifying a molecule that binds to high bone mass gene or its
                                                                                                                                                                                                                                                                                                                                                                                    Human HBM STS marker forward primer #94.
                                                                                                                                                                                                                                                                                                                                                                                                                                      02-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         corresponding wild type gene.
     11-MAY-2001; 2001US-0290071P
                                                    13-MAY-2002; 2002WO-US014876
                                                                                                                                                           WO200292764-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACC45608;
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                                                                                                                                                                                                        Homo sapiens
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ABZ22886
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Olig
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KW Phos
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KW Est
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to novel transgenic animals expressing the high CC bone mass (HBM) gene, expressing the corresponding wild type HBM gene, CC comprising an alteration of the gene encoding LRP5 or LRP6, or expressing CC an LRP5 that is modulated by an altered gene control sequence introduced CC by homologous or non-homologous recombination. The transgenic animals are CC for the study of bone density modulation or bone mass modulation. The CC invention has osteopathic and cytostatic activity. The polynucleotides of the invention may have a use in gene therapy. The transgenic animals and CC mucleic acids are for the study of bone density modulation, where the species in more than one parameter selected from bone density, bone CC strength, trabecular number, bone size, or bone tissue connectivity. The transgenic animals, nucleic acids and methods are useful for identifying CC molecules involved in bone development, and for developing pharmaceutical CC compositions, which may be employed for treating or preventing bone CC neoplasms of the bone. The transgenic animals and nucleic acids are also CC useful in methods for diagnosing diseases involved in bone development, cor characterised by reduced bone density or mass. The present sequence is useful in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                               RESULT 451
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Best Local S
Matches 17
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01-FEB-2002; 2002US-0353058P.
04-MAR-2002; 2002US-0361293P.
                                                                                                                                                                                                                                          Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New transgenic animals (e.g. mice), useful as models for studying bone density modulation, developing drugs for treating or preventing bone diseases (e.g. osteoporosis), or diagnosing diseases characterized by reduced bone density.
                                                                                                                                     Synthetic
                                                                                                                                                                                       cytostatic;
                                                                                                                                                                                                         Phosphorothicate; locked nucleic acid; LNA; immunostimulatory;
                                                                                                                                                                                                                                                                           07-APR-2003
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                                                                                                                                                                                                                                                                                                                                               ABZ22886 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 4 A; 7 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure, Page 55; 603pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Babij P,
   WO2002102825-A2.
                                                                                  modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (AMHP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          3924 GTTCCTGGGTGAGATCAA 3941
                                                                                                                                                                                                                                                                                                                                                                                                                                   19 GTTCCTGGGCGAGATCAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENOME.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bex FJ,
                                                                                                                                                                                       antimicrobial;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 THERAPEUTICS CORP
                                                                                                                                                                                                                                             /*tag=
/mod_ba
                                                                                                    Location/Qualifiers
                                   /note= "5'-terminally modified by fluorescein"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Yaworsky PJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
                                                  _base= OTHER
                                                                                                                                                                                       gene therapy; pathogenic infection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 16.4; DB 1
Pred. No. 6.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bodine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                           cancer;
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0;

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RESULT 452
ADB98306/c
ID ADB983
XX ADB983
XX ADB983
XX O4-DEC
XX Sequen
XX OSteop
KW Osteop
KW bone m
XX bone m
XX D7 HOMO &
XX HOMO &
XX HOMO &
XX I1-MA1
PR 11-MA1
PR 11-MA1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC comprising an oligonuclectide having at least one locked nucleic acid (C based on a functional molecy. Also described: (1) a complex (II) (C comprising (I) and a DNA sequence having a complementary sequence to the cligonuclectide, and encoding a gene under the control of a promoter; (2) (C a pharmaceutical composition (III) comprising (II) and a carrier or C diluent; (3) a device loaded with (III); and (4) an oligonuclectide (IV) (C comprising a first region comprising an oligonuclectide sequence having (C at least one LNA, and a second region comprising an immunostimulatory (C oligonuclectide region containing at least one unmethylated CG di-cuclectide motif. (I) has cytostatic and antimicrobial activities, and (C can be used in gene therapy. (I) and (II) can be used in medicine, and in (C can be used infections or cancer. (I) is useful for the preparation of (C pathogenic infections or cancer. (I) is useful for the preparation of (C (III), by hybridising (I) with a plasmid capable of expressing a gene (C complex with a pharmaceutical carrier. The present sequence represents an (C coligonuclectide, which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel locked nucleic acid conjugate useful in manufacturing a medicament for treating or preventing pathogenic infections or cancer, has an oligonucleotide having locked nucleic acid based on a functional moiety.
 11-MAY-2001; 2001US-0290071P.
17-MAY-2001; 2001US-0291311P.
01-FEB-2002; 2002US-0353058P.
                                                                                                                                                                                                                        Osteopathic; Gene therapy; High Bone Mass; HBM; LRP5; Zmax1; LRP6; bone mass modulation; osteoporosis; STS; sequence tagged site; ds.
                                                                                                                                                                                                                                                                            Sequence tagged site #187 used to prepare Zmax1 (LRP5) gene region map
                                                                                                                                                                                                                                                                                                                  04-DEC-2003
                                                                          13-MAY-2002; 2002WO-US014877.
                                                                                                                                                   WO200292000-A2
                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                         ADB98306
                                                                                                                                                                                                                                                                                                                                                                                           ADB98306 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 19 BP; 9 A; 0 C; 10 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention describes a locked nucleic acid (LNA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 5; Page 36; 101pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-JUN-2002; 2002WO-GB002728
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1181 GAGAAAGAGAGAGAGAGA 1198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GAGAGAGAGAGAGAGA 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.4;
Pred. No. 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4; DB 1; Length 19;
6.3e+02;
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RESULT 453
AXA95521
ID AAX9555
XX AX9555
XX AX955
XX PCR px
XX Respir
KW Respir
KW Respir
KW Incutra
XX Synthe
OS Chlamy
YX W09927
XX W09927
XX W09927
XX GEST
XX GEST
XX GEST
XX GESOM
PR 04-NON
PR
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to High Bone Mass (HBM), LRP5 (Zmax1) and LRP6 mutants, which results in a HBM-like phenotype when expressed in a cell. The HBM-like phenotype results in bone mass modulation and/or lipid level modulation. The invention is useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject suffering from e.g. osteoporosis. The present sequence is a Sequence Tagged Site (STS) osteoporosis, which was used to prepare a physical map of the Zmax1 (LRP5) gene marker, which was used to prepare a physical map of the Zmax1 (LRP5) gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19
                                                                                 Genome sequence of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                     21-NOV-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX95521;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid comprising a mutation in LRP5 or LRP6, useful for diagnosing a HBM-like phenotype in a subject and for preparing a composition for modulating bone mass and/or lipid levels in a subject
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Allen K,
                                                                                                                                      WPI; 1999-357842/30.
                                                                                                                                                                                                                                                                                                                                                                                   20-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                        03-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO9927105-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chlamydophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neutralising epitope; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer used to amplify an ORF of Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX95521 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 62; 629pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-129214/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-MAR-2002; 2002US-0361293P
                                                                                                                                                                                                                                             (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (AMHP ) WYETH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
les 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3924 GTTCCTGGGTGAGATCAA 3941
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anisowicz
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                pneumoniae.
                                                                                                                                                                                                                                                                                                     97FR-00014673.
98US-0107078P.
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                                                                                                                                                                                                                                                                                                                                                                                   98WO-IB001890.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
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Pred. No. 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       erythema nodosum; pharyngitis; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Morales A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 19;
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Page 1754; Disclosure; 1912pp; English.

ត្តទុក្ខភព្គក្ខិត្តក្នុង ស្ត្រីក្រុង ស្ត្រីក្រុង ស្ត្រីក្រុង ស្ត្រីក្រុង ស្ត្រីក្រុង ស្ត្រីក្រុង ស្ត្រីក្រុង ស្ត្រីក្រុង ស្ត្រីក្រុង ស្

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RESULT 454
AAAH777C
ID AAAH7777
XX AAH7777
XX AAH7777
XX Thromb
KW Thromb
KW Thromb
KW Dosin
XX Homo 8
XX Homo 8
XX E-PUI
PR 29-JUI
PR 29-JUI
PR 29-JUI
PR 29-JUI
PR 11-JAN
PR 1
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ota T, Ishii S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, crythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAY34584- AAY35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising
                                                                             PCR primers AAH77766-67 were used to amplify a cDNA fragment encoding thrombospondin 1-like (TSP1-like) protein. The cDNA sequence encoding human TSP1-like protein was isolated from a human 10 week-aged foetal tissue cDNA library. The TSP1-like protein may be involved in intracellular vesicle transport. Secretion of the TSP1-like protein is reduced in brain tumours. It can be used in the screening of target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               767/c
AAH77767 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JUL-1999;
27-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH77767;
                                                   compounds,
                                                                                                                                                                                                                                                                           Example 10; Page 34;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-JAN-2000;
17-FEB-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-JUL-2000; 2000WO-JP005068
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-FEB-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Thrombospondin 1-like protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       13-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (HELI-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1620 CTTCAGCTGCAGAGAGCT 1637
                                                                                                                                                                                                                                                                                                                            encoding thrombospondin-like protein, t, useful for diagnosis and treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                     2001-541222/60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                tumour; PCR primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isogai T, Nishikawa T,
, Sugiyama T, Wakamatsu
Kanzaki K, Inoue Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ç
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kanzaki K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTTCATCTGCAGAGAGCT 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C. pneumoniae
                                                      and is useful for diagnosis, prediction and treatment of brain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    99JP-00300253.
99US-0159590P.
2000JP-00118776.
2000US-0183322P.
2000JP-00183767.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
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                                                                                                                                                                                                                                                                           105pp; Japanese
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Pred. No. 6.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TSP1-like protein; vesicle transport;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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A, Nagai K,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Saito K, Ya
, Otsuki T,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Yamamoto
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                                                                                                       Matches
                                                                                                                                                                                                                                                         The invention comprises antisense oligonucleotides that are capable of inhibiting expression of the human clusterin gene. The antisense oligonucleotides of the invention are useful for inhibiting the expression of clusterin in cells. The antisense oligonucleotides are useful for treating an animal with a disease or condition associated clusterin (e.g. hypercholesterolaemia; cardiovascular disorders; hyperproliferative disorders; and hyperlipidemic disorders). The presents a clusterin antisense oligonucleotide of the invention. NOTE: The present DNA sequence has a phosphorothicate back and also contains 2'-O-methoxyethyl wings
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel antisense compound targeted to nucleic acid molecule encoding clusterin, useful for treating animal having disease associated with clusterin such as hyperlipidemic disorder, cardiovascular disorder.
                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2002-404805/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10-SEP-2001; 2001WO-US028235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypercholesterolaemia; cardiovascular disorder; ss; hyperproliferative disorder; hyperlipidemic disorder; phosphorothioate backbone; 2'-O-methoxyethyl wing.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human clusterin inhibiting antisense oligonucleotide 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-AUG-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-SEP-2000; 2000US-00659791.
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 84; 125pp;
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                                                     CTGGGTGTCCTGGCTGGC 2099
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                                                                                                                                                                                                               BP; 4 A; 8 C; 6 G; 2 T; 0 U; 0 Other;
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                                                                                                          Conservative
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                                                                                                                             94.48;
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                                                                                                                                                         0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 English.
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Pred. No. 6.
                                                                                                                                Score 16.4; DB 1
Pred. No. 6.4e+02
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                                                                                                                                                                                               nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antiasthmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of abquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nyce Ji
Miller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a novel pharmaceutical composition, which has first active agent comprising an oligonuclectide antisense to the initiation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; bypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-APR-2001; 2001US-0286137P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              junctions of genes encoding a polypeptide associated with lung and/or
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                                                                                          Similarity
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, Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQ ID NO 1318; 872pp; English
                                                                                                                                             BP; 1 A; 6 C; 7 G; 6 T; 0 U; 0 Other;
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L, Shahabuddin
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                                                                                        6.4e+02
                                                                                                         DB 1; Length 20
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Composition of larget polypeptide associated with lung or reduction has antiallergic, and for useful composition or cancer and for use of the invention has antiallergic, antiinflammatory, altered to a mammal. The coligonucleotides are derived from a gene encoding or regulating construction or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating construction of a target polypeptide associated with lung airway or lung construction or cancer and can be anti-sense to the corresponding mRNA. Constructions for adding a carrier and for use of the corresponding mRNA. Constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, an algesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition is useful for lungs and or target polypeptide present in the lungs. The composition and/or bronchoconstriction are associated with a disease or condition such as pulmonary vasoconstriction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory disease; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis;
                                                                                                          inflammation, allergies, asthma, impeded respiration, respiratory distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary hypertension, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the present in the target RNA serves to prevent the breakdown of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                            the oligonucleotides into products that free adenosine into e.g., lung, brain, heart, kidney, etc, tissue environment ar prevent any unwanted effects due to it
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes a novel composition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 15; SEQ ID NO 1318; 763pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-APR-2001; 2001US-0286036P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pulmonary transplantation rejection; ss; primer.
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, Tang L,
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L, Shahabuddin S;
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RESULT 458
ADJ53373/G
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                                                                                                         Query Match
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Matches 17
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                                                                                                                                                                                                                                                                                        The invention relates to an antisense oligonucleotide targeted to a nucleic acid encoding the human G protein-coupled receptor 6 (GPCR-6), which specifically hybridises with the nucleic acid encoding the GPCR-6 and inhibits expression of the GPCR-6. The antisense oligonucleotide comprises at least one modified internucleoside linkage, i.e. a phosphorothioate linkage, at least one modified sugar molety, preferably a 2'-O-methoxyethyl sugar molety, or at least one modified nucleobase comprising a 5-methylycycosine. The antisense oligonucleotides are useful for inhibiting expression of the GPCR-6 and in preparation of a composition for treating a disease or condition associated with GPCR-6, e.g., a metabolic disorder, aberrant signal transduction in brain tissue, a neuronal, motor, sensory, psychiatric or behavioural disorder or drug or chemical addiction. This sequence represents an antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel antisense compound targeted to nucleic acids encoding G protein-coupled receptor 6 (GPCR-6), useful for treating animal having disease associated with GPCR-6 e.g. metabolic, neuronal, motor, sensory or behavioral disorders.
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aberrant signal transduction; brain tissue; neuronal disorder;
motor disorder; sensory disorder; psychiatric disorder;
                                                                                                                                                                                                                   Sequence
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                                                                                                                                   Similarity
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                                                                                                                                                          Score 16.4;
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Pred. No. 6.4e+02;
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RESULT: 459
ADJ53441
ID ADJ5344
XX ADJ534
XX A
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                                                                                                                                                                                   Query Match
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Matches 17
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Human; G protein-coupled receptor 6; 
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aberrant signal transduction; brain tissue; neuronal disorder;
motor disorder; sensory disorder; psychiatric disorder;
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                                                                                                                                                                                        Conservative
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Pred. No. 6.4e+02;
                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                       ۲.
                                                                                                                                                                                                                                                                             Length 20;
                                                                                                                                                                                        Indels
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                                                                                                                                                                                        Gaps
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AAQ20035/c ID AAQ200 XX AC AAQ200

AAQ20035 standard; DNA; 21

BP

AAQ20035;

RESULT 460

밁 5

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The sequence is designed to target the Human tumour necrosis beginning at nucleotide 1137 and to covalently cross-link to
                                                                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Key
modified_base
                     Example 4; Page 25; 42pp; English.
                                             New sequence-specific non-photo-activated crosslinking agents -
the major groove of duplex DNA and are esp. useful for treating
                                                                                      Matteucci MD,
                                                                                                                     25-MAY-1990;
14-JAN-1991;
                                                                                                                                            25-MAY-1990;
                                                                                                                                                             12-DEC-1991.
                                                                                                                                                                            WO9118997-A.
                                                                                                                                                                                                                  modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               deoxyribonucleic acid; major groove; ethanoamino group;
aziridinylcytosine; cross-linking group; tumour necrosis factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cross-linking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-APR-1992
                                                                       WPI; 1992-007480/01.
                                      infections e.g. HIV.
                                                                                                      (GILE-) GILEAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       oligomer 216 for targetting human
                                                                                     Krawczyk S;
                                                                                                      SCIE INC.
                                                                                                                     90US-00529346.
91US-00640654.
                                                                                                                                             90US-00529346
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/mod_base=
                                                                                                                                                                                                                                            /*tag=
/mod_ba
17
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/mod_base=
                                                                                                                                                                                       /mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine"
                                                                                                                                                                                                                                                                                                 /mod
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                                                                                                                                                                                                                                                                                                                                                                                                              /mod
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                                                                                                                                                                                                                                                                                                                                                                      /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "N-methyl-8-oxo-2'-deoxyadenine"
                                                                                                                                                                                                                                                                                                                               *tag=
                                                                                                                                                                                                                                                                                                                                              mod_base= m5c
                                                                                                                                                                                                                                                                                                                                                                                                    note= "N-methyl-8-oxo-2'-deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note= "N4N4-ethanocytosine"
                                                                                                                                                                                                          tag=
                                                                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                base= OTHER
== "N-methyl-8-oxo-2'-deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           _base= OTHER
                                                                                                                                                                                                                                                  base=
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factor
it via the
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RESULT 461
AAQ30385;
ID AAQ30385;
XX
AC AAQ30385;
XX
DT 25-MAR-2003 (revised)
DT 07-DEC-1992 (first entry)
XX
DE Oligomer TNP216 for forming t
XX
Tumour necrosis factor; herpe
XX
XX
Tumour necrosis factor; herpe
XX
XX
FH Key
FT modified_base 1 Location/Qual
FT modified_base 2 /*tag= a
FT //note= "OTHE!
FT modified_base 3 /*tag= c
FT //note= "OTHE!
FT modified_base 4 /*tag= d
FT //note= "OTHE!
FT modified_base 7 //note= "OTHE!
FT modified_base 7 //nod_base= o/
FT //nod_base= m
FT modified_base 1 /*tag= d
FT //nod_base= m
FT modified_base 1 /*tag= f
FT modified_base 1 /*tag= i
FT modified_base 1 /*tag= j
FT modified_base 1 /*tag= i
FT modified_base 1 /*tag= j
FT modified_base 2 /*tag= k
FT modified_base 2 /*tag= k
FT /*tag
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Best Local Similarity 94.4%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N4N4-ethanocytosine group. See also AAQ20031-Q20038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tumour necrosis factor; herpes simplex; AIDS; modified; HIV; RSV; HPV; malignancy; hepatitis; inflammation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligomer TNF216 for forming triplex with HUMTNFAA target duplex.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                  /mod_base= OTHER
/note= "OTHER= N6 methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= OTHER
/note= "OTHER= N6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "OTHER= N4
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== "OTHER= N6
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= "OTHER= N6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              methyl-8-oxo-2'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   methyl-8-oxo-2' deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ethanocytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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RESULT 462
AAQ30382/c
ID AAQ303
XX AAQ303
XX AC AAQ303
XX DT 07-DEC
XX Oligom
XX Tumoux
XW Tumoux
XW Tumoux
XW Tumouifi
FT modifi
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human tumour necrosis factor beginning at nucleotide 1137 contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV, HER, HIV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-NOV-1990;
18-JAN-1991;
08-APR-1991;
17-APR-1991;
17-APR-1991;
17-APR-1991;
                                                                                                                                Key
modified_base
                                                                                                                                                                                                    Tumour necrosis factor; herpes simplex; AIDS; modified; HIV; RSV; HPV; mallgnancy; hepatitis; inflammation; ss.
                                                                                                                                                                                                                                           Oligomer TNF213
                                                                                                                                                                                                                                                                     25-MAR-2003
07-DEC-1992
                                                                                                                                                                                                                                                                                                                                     AAQ30382 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New oligomers contg. modified bases - doublet in a DNA duplex, for treating herpes malignancy and inflammation.
                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                            AAQ30382;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 4 A; 7 C; 0
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                            modified_base
                                                                                modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GILE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                 1184 AAAGAGAGAGAGAAAT 1201
                                                                                                                                                                                                                                                                                                                                                                                                                                                           17;
                                                                                                                                                                                                                                                                                                                                                                                                      20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GILEAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ₽,
                                                                                                                                                                                                                                                                                                                                                                                                      AAAGAGAGAGAGAATT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Krawczyk S,
                                                                                                                                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90US-00617907.
91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
91US-00766733.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SCI INC.
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                                                     /*tag= b
/mod_base= (
                                                                                                                                                                                                                                           for forming triplex with HUMTNFAA target duplex.
/*tag= c
/mod_base= OTHER
                                                                                              /mod_ba
                                                                                                                                                 Location/Qualifiers
                                           /note=
                                                                                                          /*tag= a
/mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
                                              "OTHER=
                                                                                                "OTHER=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matteucci MD,
                                             OTHER
HER= N6
                                                                                       OTHER
HER= N6 methyl-8-oxo-2'
                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 16.4; DB 1;
Pred. No. 6.4e+02;
0; Mismatches 1
                                             methyl-8-oxo-2'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      which form a triplex with G-C and diagnosing HIV, hepatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Milligan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                             deoxyadenine"
                                                                                                deoxyadenine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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17-APR-1991;
17-APR-1991;
17-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-NOV-1990;
18-JAN-1991;
08-APR-1991;
The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human tumour necrosis factor beginning at nucleotide 1137 contg. a purine rich sequence concd. on one strand of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. HPV; HBR; HIV, hepatitits B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                     New oligomers contg. modified bases - which form a triplex with doublet in a DNA duplex, for treating and diagnosing HIV, hepatiherpes malignancy and inflammation.
                                                                                                                                                                                                                                 Claim 12; Page 70; 77pp; English.
                                                                                                                                                                                                                                                                                                                                                                                       Froehler B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-NOV-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-JUN-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                               (GILE-)
                                                                                                                                                                                                                                                                                                                                                                                                                             GILEAD
                                                                                                                                                                                                                                                                                                                                                                                         Krawczyk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90US-00617907
91US-00643382
91US-00683420
91US-00686544
91US-00686546
91US-00686547
91US-00686547
                                                                                                                                                                                                                                                                                                                                                                                                                               SCI INC.
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/mod_base= OTHER
- "OTHER= N6
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/mod_base=
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/mod_ba
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                       Matteucci MD,
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                                                                                                                                                                                                                                                                                                                                                                                         Milligan
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                                                                                                                                                                                                                                                                                        ex with G-C
, hepatitis,
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1184 AAAGAGAGAGAGAAAT 1201

Matches Query Match

Local

17; Conserv

Conservative

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0.3%;

Score 16.4; DB 1 Pred. No. 6.4e+02 Mismatches

DB 1;

Length 21;

Indels

0

Gaps

٥,

Sequence 21 BP; 5 A; 6 C; 0 G; 10 T; 0 U; 0 Other;

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RESULT 463
AAH62152
ID AAH6211
AC AAH621
XX AAH621
XX O9-SE
DT 09-SE
DT 12-SE
XX Singl
XX Homo
OS Unid
XX Homo
OS Unid
XX Homo
OS Unid
XX FT Vari
FT Vari
FT Vari
FT YX
PD 31-
XX AY
PD 31-
XX AY
PD 17-
XX AY
PD 24-
XX AY
PD 17-
XX AY
PD 24-
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                                                                                                                                             Query Match
Best Local :
                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                            DNA sequences AAH62100 - AAH62688 represent segments of human genes which contain single nucleotide polymorphisms (SNPs). A method is included in the invention for analysing a nucleic acid sample, which consists of determining the base occupying any one of the polymorphic sites given in the SNP containing sequences. The nucleotide sequences can be used in the diagnosis or monitoring of diseases, such as cancer, inflammation, heart diseases, diseases of the cardiovascular system, and infection by microorganisms. The oligonucleotides are also useful in the manufacture of a medicament for the treatment or prophylaxis of the diseases, and as a pharmaceutical. SNP containing oligonucleotides are useful in applications such as phenotype correlation, forensics, paternity testing, medicine and genetic analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New nucleic acid segments of the human genome, particularly from genes including polymorphic sites, for phenotype correlation, forensics, paternity testing, medicine and genetic analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism; SNP; human; cancer; inflammation; heart disease; paternity testing; forensic science; ds.
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12-SEP-2001
                                                                                                                                                                                                                                Sequence 21 BP; 3 A; 8 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                 Revised
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 34; 80pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2001-367705/38.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cargill M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-NOV-2000; 2000WO-US031639
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Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WHED ) WHITEHEAD INST BIOMEDICAL RES.
                                                       1583 GCCAGCTGTATGGGGACC 1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
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                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                              record issued on 09-SEP-2004 : Correction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ireland JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /*tag= a
/standard_name= "single nucleotide polymorphism"
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                                                                                                                                             0.3%;
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                                                                                                              <u>,</u>
                                                                                                                                                                         Score 16.4;
                                                                                                                                             Pred.
                                                                                                                  Mismatches
                                                                                                                                             No.
                                                                                                                                             6.4e+02
                                                                                                                                                                         DB 1; Length 21;
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                                                                                                                                                                                                                                                                                              Feature Table Key
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RESULT 464

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RESULT 465
ADJ13846/c
ID ADJ138
XX
AC ADJ138
XX
DT 20-MAY
XX
DE Human
XX
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                         modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                 Human DNA probe used to immobilise CpG methylated DNA SeqID 973
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human DNA probe used to immobilise CpG methylated DNA SeqID 828.
                                                                                                                ADJ13846 standard;
                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 2 A; 13 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; SEQ ID NO 828; 210pp; English.
                                                 20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-2001; 2001US-0301370P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-2002; 2002US-00184085
                                                                                                                                                                                                                                 2436 GGATGAGAAGGGGAGAGG 2453
                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ss; chemical modification; methylation; array; CpG island
                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                  GGATGAGAGGGGGAGAGG 3
                                                                                                                                                                                                                                                                 Conservative
                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             p16; human; H69; H1618.
                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                 94.4%;
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                                                                                                                  21
                                                                                                                  BP
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                                                                                                                                                                                                                                                                                 Score 16.4; DB 1; Length 21; Pred. No. 6.4e+02;
                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Balog
                                                                                                                                                                                                                                                                   Indels
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DNA

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RESULT 466
ADJ13700/C
ID ADJ137
XX
ADJ1370/C
AC ADJ137
XX
DT 20-MAY
XX
DB Human
XX
DB Human
XX
DB Humour
XX
C
DB Homo B
XX
C
DS Homo B
XX
DS HOM
DS HO
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Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        particular, this invention reters to analysing the mountained from two a region of the promoter for the tumour suppressor gene p16 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Analysis of chemical modification of DNA involves obtaining sample of to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Garner HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probe; ss; chemical modification; methylation; array; CpG island;
tumour suppressor; p16; human; H69; H1618.
                                   14-AUG-2003
                                                                                        US2003152950-A1
                                                                                                                                                                                                                                                                                                                                                20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADJ13700 etandard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel method for analysing chemically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-874843/81.
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                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                        DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19
                                                                                                                                                                                                     se; chemical modification; methylation; array; CpG island, suppressor; p16; human; H69; H1618.
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MINNA J D.
LUEBKE K J.
BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       suppressor; p16; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGATGAGATGGGGAGAGG 2
                                                                                                                                                                                                                                                                                        probe used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                  to immobilise CpG methylated DNA SeqID 827.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         210pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 16.4; DB 1; Length 21; Pred. No. 6.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 6 T; 0 U; 0 Other;
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RESULT 467
ADJ13810/c
ID ADJ138
XX ADJ138
XX ADJ138
XX ADJ138
XX Probe;
XW probe;
XW tumour
XX Homo 8
XX Homo 8
XX 12003
PD 14-AUG
XX 27-JUN
XX 27-JUN
XX PA (GARN/PA (LUEB)
PA (BALO/XX
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Analysis of chemical modification of DNA involves obtaining sample to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA
               (MINN/)
(LUEB/)
(BALO/)
                                                                                                                                                    14-AUG-2003
                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                             tumour
                                                                                                                                                                                                                                                                                                                    20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                    ADJ13810;
                                                                                                                                                                                                                                                                                                                                                                                ADJ13810 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 2 A; 13 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; SEQ ID NO 827; 210pp; English.
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                                                                                           27-JUN-2001; 2001US-0301370P
                                                                                                                      27-JUN-2002; 2002US-00184085
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention.
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(MINN/) MINNA J D.
(LUEB/) LUEBKE K J.
                                                           (GARN/)
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                                                                                                                                                                                                                                                                                      DNA probe used to immobilise CpG methylated DNA SeqID 937.
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               LUBBKE K J.
BALOG R P.
                                                                                                                                                                                                                                                          ss; chemical modification; methylation;
                                                                                                                                                                                                                                          suppressor; p16; human; H69; H1618.
                                                           GARNER H R.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            GGATGAGAGGGGGAGAGG
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                                                                                                                                                                                                                                                                                                                      (first entry)
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Pred. No. 6.4e+02;
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                                                                                                                                                                                                                                                          array; CpG island
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RESULT 468
ADJ13739/c
ID ADJ137
XX ADJ137
XX ADJ137
XX Probe;
KW probe;
KW tumoux
XX Homo E
XX US2003
XX Homo E
XX US2003
XX Homo E
XX US2003
XX US2003
XX Homo E
XX GARN,
PA (MINN,
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PA (HALO,
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XX GARN)
PA (BALO,
XX GARN)
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Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                Analysis of chemical modification of DNA involves obtaining sample of to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADJ13739 standard; DNA; 21 BP
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                                                                                                                                                           WPI; 2003-874843/81.
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                                                                                                                                                                                                                                                                                                                                        (GARN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-AUG-2003
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                                                                                                                                                                                                                                                                                                                (MINN/
                                                                                                                                                                                                                                                                (/OTAB)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2436 GGATGAGAAGGGGAGAGG 2453
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                                                                                                                                                                                                                                                           ) GARNER H R.
) MINNA J D.
) LUEBKE K J.
) BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; chemical modification; methylation; array; CpG island.
suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               probe used to immobilise CpG methylated DNA SeqID 866.
        SEQ
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                                                                                                                                                                                                           Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 2 A; 12 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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210pp;
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Pred. No. 6.4e+02;
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RESULT 469
ADJ13039/c
ID ADJ130
AC ADJ130
AC ADJ130
AC ADJ130
AX Probe;
KW Probe;
KW Cumoux
XX Homo E
AX US2003
AX CARN,
PA (ALAN,
PA (MINN,
PA (BALO,
XX (B
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Matches 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human DNA probe used to immobilise CpG methylated DNA SeqID 166.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Garner HR, Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probe; ss; chemical modification; methylation;
tumour suppressor; p16; human; H69; H1618.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (BALO/)
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LUEBKE K J.
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Pred. No. 6.
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method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequence.

DNA sequences

This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput

Example 1; SEQ ID NO 166; 210pp; English

Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.

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RESULT 470
ADJ13774/c
ID ADJ1377
XX ADJ137
XX ADJ137
XX Probe;
XX Probe;
XX Probe;
XX US2003
XX US2003
XX US2003
XX US2003
XX (GARN)
PA (MINN)
PA (MINN)
PA (MINN)
PA (BALO)
XX WPI;
XX WPI;
XX WPI;
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XX C
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Best Local S
Matches 17
                                                                               This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Analysis of chemical modification of DNA involves obtaining sample of to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pi6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-874843/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GARN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-2002; 2002US-00184085
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 that comprise the modifications of interest such as a CpG island. In
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (BALO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2436 GGATGAGAAGGGGAGAGG 2453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GARNER H R.

MINNA J D.

LUEBKE K J.

BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; chemical modification; methylation; array; CpG island;
suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Minna JD,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        901;
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Pred. No. 6.
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Sequence

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Other,

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Query Match Best Local S Matches 17

Similarity

94.4%;

0.3%;

Score 16.4; DB 1 Pred. No. 6.4e+02

DB 1;

Length

21

Indels

0

Gaps

0,

Conservative

0;

Mismatches

GATGAGAAGGGGAGAGGT 2454

Sequence 21

BP; 4 A;

12 C; 0 G; 5 T; 0 U; 0 Other;

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Best Local 9
                                   This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pil from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                             Analysis of chemical modification of DNA involves obtaining sample to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA
                                                                                                                                                                                                                                     Example 1;
                                                                                                                                                                                                                                                                                                                       WPI; 2003-874843/81.
                                                                                                                                                                                                                                                                                                                                                  Garner HR, Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                               27-JUN-2001; 2001US-0301370P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 probe; ss; chemical modification; methylation; array; CpG island;
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                                                                                                                                                                                                                                                                                                                                                                             (BALO/)
                                                                                                                                                                                                                                                                                                                                                                                                      (GARN/) GARNER H R.
(MINN/) MINNA J D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   suppressor; p16; human; H69; H1618
                                                                                                                                                                                                                                                                                                                                                                            LUEBKE K J.
BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                     SEQ ID NO 234; 210pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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Pred. No. 6
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                                                                                                                                                                                                                                                                                              DNA
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ARGSULT 473
AAQG7954/C
ID AAQG79
XX AAQG79
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XX DAQG79
XX DAQG79
XX CAQG79
XX CAQG79
XX CQ44 P
XX CQ44;
XX PCR; ax
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ABT21572
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel method for determining racing potential of a horse. The method comprises measuring: whether grandpaternal DNA is over-represented in the genome of the horse; or in the case where one of the grandmothers was selected for breeding on the basis of racing performance, whether grandmaternal DNA from the selected grandmother is over-represented in the genome of the horse which indicates that the horse has good racing potential. The method of the invention is useful for determining the racing potential of a horse or for obtaining a progeny horse with good racing potential. This polymucleotide sequence represents a PCR primer used in the detection method of over-representation of DNA from male grandparents of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Determining the racing potential of a horse comprises measuring whether grandpaternal or grandmaternal DNA from the selected grandmother DNA is over-represented in the genome of the horse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Racing potential; horse; grandpaternal DNA; over-represented; breeding; grandmother; performance; progeny horse; PCR; primer; ss.
                     CD44; exon 6; cell surface protein; primer; polymerase chain reaction; PCR; amplification; tumor; imaging; therapy; ss.
                                                                                            CD44 PCR primer
                                                                                                                                       25-MAR-2003
08-DEC-1994
                                                                                                                                                                                                            AAQ67954;
                                                                                                                                                                                                                                                       AAQ67954 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 22 BP; 1 A; 8 C; 2 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 2; Page 25; 49pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Multiplex group PCR primer #319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16-APR-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Binns MM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-MAY-2001; 2001GB-00011886
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200292851-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2003-129314/12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-MAY-2002; 2002WO-GB002273
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ANIM-) ANIMAL HEALTH TRUST.
(BRHO-) BRITISH HORSERACING BOARD
                                                                                                                                                                                                                                                                                                                                                                                                               4870 TCTCAGTTTCTTCTCTG 4887
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                                                                                                                                       (revised)
(first en
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                                                                                                                                       entry)
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                                                                                                                                                                                                                                                       BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 6.40
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 16.4; DB 1;
Pred. No. 6.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1; Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 474
AAQ55904/c
ID AAQ559
XX AAQ559
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XX PCR pr
XX PCR; p
XX PCR; p
XX BAGNC
XX Synthe
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Best Local :
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16-DEC-1992;
20-JUL-1993;
                                                                                                                                                                                    21-JUL-1992;
20-NOV-1992;
16-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tarin D,
  Method for diagnosing neoplasia,
                                                                                                                                                                                                                                                                                                                                                                      WO9402633-A1
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25-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI;
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                                                                                          Barin D,
                                                                                                                                                                                                                                                                              20-ЛИТ-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ55904;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-JUN-1994.
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SXCCCCCXXXXIII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The primers given in AAQ67948-58 are used for cancer diagnosis assay. The primers given in AAQ67948-51 anneal to the standard CD44, and those given in AAQ67952-58 anneal to exon sequences on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                        PCR; polymerase chain reaction; CD44; cancer; bladder cancer; detection; diagnosis; carcinoma; solid tumour; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 5 A; 6 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                        WPI; 1994-048890/06.
                                                                                                                                                                                                                                                                                                                                                                                        AAQ55904 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 25; 73pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antibody specific for peptide of exon 6 of CD44 - also exon 6 fragments and fusion protein, useful for diagnosis, treatment and
                                                                          (ISIS-) ISIS INNOVATION LTD
                                                                                                                                                                                                                                                                                             PCR primer for amplifying region of CD44 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3255 CCAGGACCTGGCCTCTGTGCT 3275
                                                Matsumura
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matsumura
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                                                                                                                                                                                                                                                                                                                         (revised)
(first entry)
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                                                                                                  92GB-00015498.
92GB-00024386.
92GB-00026165.
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92GB-00026165
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Pred. No. 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 21;
utilises unequal
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(Updated
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imaging
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RESULT 475
AAQ75682/c
ID AAQ756
XX AAQ76

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ៥៩៩៩ភព្វាក្សាក្រុក្សព្រះក្រុក្សព្រះក្រុក្ស
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11 primers (AAQ55898-Q55908) were used in the amplification of a CD44 gene region and in the subsequent detection of the amplification products. cDNA derived from urine of 90 patients (44 with biopsy proven bladder cancer, 46 from normal volunteers and patients with neoplastic inflammation of the bladder) was divided equally into two tubes, one for PCR with primers B1 (AAQ55902) and B5 (AAQ55906) to amplify the transcript for diagnosis, the other for PCR with P1 (AAQ55998) and P4 (AAQ55901) to amplify the standard form of CD44. When all exons were expressed (bladder cancer), B1 and B4 produced a 735 bp band. There was
A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         no band in tracke containing cDNA from normal patients or those with bladder inflammation. A 482 bb band in all cases with P1 and P4 indicated that diagnostically significant differences between urine from patients with bladder cancer and controls was due to alternative splicing of the CD44 gene. The method allows diagnosis of neoplasia based on overexpression of the CD44 exon. Solid tumours, especially carcinomas, can be accurately detected. This primer is complementary to a sequence contained within exon 11 of the CD44 gene. (Updated on 25-MAR-2003 to correct PN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ75682;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ75682 standard;
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                                                                                                                                                                                                                                                                                                            Analysis of cDNA and gene exp
by digestion with restriction
                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example
                                                                                                                                                                                                                                                  Disclosure; Page 7; 11pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        splicing of CD44 gene in cancer patients compared to controls.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3255 CCAGGACCTGGCCTCTGTGCT 3275
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCAGGAACTGTCCTCTGGGCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene expression; reverse transcription; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 17; 42pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                            expression -
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Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                               enzymes.
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                                                                                                                                                                                                                                                                                                                                        amplification
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
   seperate lanes. The
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RESULT 477 AAQ90391

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AAQ90391 st. AAQ90391; 08-JAN-1996

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                                                        Query Match
Best Local 8
                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local :
                                                                                                                  A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (CBNESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21 BP; 1 A; 1 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                              Sequence 21
                                                                                                                                                                                                                                 Disclosure; Page 6; llpp; Japanese.
                                                                                                                                                                                                                                                        Analysis of cDNA and gene expression by digestion with restriction enzymes
                                                                                                                                                                                                                                                                                                                                              16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                       16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reverse transcription primer used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAQ75645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ75645 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       method
                                                                                                                                                                                                                                                                                                                      (NITE ) NIPPON TELEGRAPH & TELEPHONE
                       5399 ATACAAAAAAGAAAAAATGAA 5419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5388 GAATTAAAAAATACAAAAA 5408
                                                                                                                                                                                                                                                                                              1995-018287/03
21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     can be used to analyse gene expression rapidly and easily
                                              18;
                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAATAAAAAAAAAAAAAA 1
ATACAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gene expression; reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       restriction enzyme; ss.
                                                                                              BP; 1 A; 0 C; 1 G; 19 T; 0 U; 0 Other;
                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                              93JP-00112515.
                                                                                                                                                                                                                                                                                                                                                                       93JP-00112515
                                                         0.3%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                              0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                         Score 16.2; DB 1
Pred. No. 6.8e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.2;
Pred. No. 6.
                                                                                                                                                                                                                                                          enzymes.
                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                       CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cDNA analysis technique.
                                                                                                                                                                                                                                                                     Ą
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6.8e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                      DB 1;
                                                                                                                                                                                                                                                                       amplification of mRNA followed
                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                       21;
                                               0
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                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                66
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RESULT 478
AAT10743
ID AAT107
XX AAT107
XX O9-SEE
XX Oligor
XX Electr
KW Electr
KW attack
KW detect
XX detect
XX Synthe
XX Synthe
XX Key
FT modif:
                                                                                                                                                                                                                                                                                                      밁
                                                                                                                                                                                                                                                                                                                            δ
                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                        The sequences represented by, AAQ90390-90401 are synthetic DNA probes containing 3' ribonucleoside termini. The sequences shown in AAQ90402-15 are synthetic DNA probes with 5' amino termini. These sequences were specific for the polymorphisms of HIA gene dQa. These sequences were used in the device of the invention. This is a self-addressable electronic device (SAED) that can be used to carry out multi-step and multiplex reactions, such as nucleic acid hybridisations. The advantages of this method are that these reactions can be carried out with complete and precise electronic control, and that the rate, specificity and sensitivity of these reactions are greatly improved at micro-locations
                                                                                  attachment detection;
                                                                                                                                         Oligonucleotide probe, CP-1.
                                                                                                                                                                        09-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 20 A; 0 C; 0 G; 0 T; 1 U; 0 Other;
                                                         Synthetic.
                                                                                                              Electronically self-addressable device; ED; electrode; current source;
                                                                                                                                                                                                                             AAT10743 standard; RNA; 21 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 40; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and bio:polymer synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New self-addressable electronic devices - used for multi-step and multiplex reactions such as DNA hybridisation(s), clinical diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-185870/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Heller MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9512808-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CP-1 (synthetic DNA probe with 3'ribonucleoside terminal #2).
              modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NANO-) NANOGEN INC
                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                                                                                                                                                                                                                                                                                   5396 AAAATACAAAAAAGAAAAAAT 5416
                                                                                                                                                                                                                                                                                                                                                         17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HLA; dQa; 3' r: hybridisation;
                                                                                                                                                                                                                                                                                                    AAAAAAAAAAAAAAAAAA 21
                                                                                     88.
                                                                                                layer; permeable; counterion; genetic typing; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7
                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ħ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          93US-00146504
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94WO-US012270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ų
                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
/*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /*tag=  a
/note= "3′ ribonucleoside terminal"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ribonucleoside; self-addressable electronic device;
                                                                                                                                                                                                                                                                                                                                                                       81.0%;
                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
                                                                                                                                                                                                                                                                                                                                                         1:
                                                                                                                                                                                                                                                                                                                                                                        Score 16.2; DB 1;
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                     Length 21;
                                                                                                                                                                                                                                                                                                                                                           Indels
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RESULT 479
AAV36491
ID AAV364
XX AAV364
AC AAV364
XX Parimex
XX Partia
KW Primex
XX Pomo E
XX PO9820
XX HOMO E
XX HOMO E
XX HOMO E
XX OOS-NON
XX O1-NON
XX O1-NON
XX OX-NON
XX O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Watches 17; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              문
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The sequences given in AAT10742-67 are synthetic oligonucleotides which are used in the construction of the electronically self-addressable device (ED) of the invention. The ED comprises a substrate, an electrode or opt. a number of electrodes supported by the substrate, a current source operatively connected to the electrode and an attachment layer adjacent to the electrode which is permeable to a counterion but not permeable to a molecule capable of insulating or binding to the electrode. The attachment layer is capable of attaching a macromolecule. The ED is used for genetic typing and comprises a number of electronically addressable locations each comprising an electrode, and a binding entity, such as one of these probes, attached to each of the locations capable of detecting the presence of a genetic sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Electronically self-addressable device e.g. nucleic acid hybridisation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-097582/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Heller MJ, Tu E, Evans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUL-1994;
  WPI; 1998-286855/25
                                                          Doxsey SJ;
                                                                                                                                                                                                                                                                                                                                           WO9820025-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Partial centrosomal protein 140; CP140; scleroderma; sclerotic disease; Primer; amplification; PCR; PCR-RFLP; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Primer R2.5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-SEP-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAV36491;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV36491 standard; cDNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO9601836-A1
                                                                                                                                                                        05-NOV-1996;
                                                                                                                                                                                                                             04-NOV-1997;
                                                                                                                                                                                                                                                                                    14-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 1; Page 60; 155pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NANO-) NANOGEN INC
                                                                                                                 (UYMA-) UNIV MASSACHUSETTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5396 AAAATACAAAAAGAAAAAAT 5416
                                                                                                                                                                                                                                                                                                                                                                                               sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAAAAAAAAAAAAAAAA 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amplifies bases 535-555 of the CP140 gene.
                                                                                                                                                                        96US-00743200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94US-00271882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95WO-US008570
                                                                                                                                                                                                                               97WO-US020520.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; 0 C; 0 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 16.2; DB 1;
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sosnowski
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ۲,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for electronic control of,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>;</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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0,

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RESULT 480
AAV57641
ID AAV576
XX AAV576
XX Exon 4
XX Epithe
XX Epithe
XX Epithe
XX Exon 4
XX Epithe
XX Homo e
XX Homo e
XX Homo e
XX I1-MAH
XX UYYA
XX UYYA
XX Determ
PT mediat
PT preser
PT preser
PT preser
PT preser
CC Sequer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 18
Sequences shown in AAV57601 to AAV57686 represent primers used for the PCR amplification of the axons of the different subunits of the human epithelial sodium channel (ENAC) gene. This is used in the method of the invention of determining the presence or absence of a mutation conferring a pathological condition mediated by altered ion transport. The method comprises analysing a nucleic acid sample, or protein sample, for the presence of a mutation in the ENAC gene, or in its encoded protein. A vector containing a nucleic acid encoding a human altered variant of the
                                                                                                                                                                                                                                                                                                 Determination of presence of mutation conferring pathological condition mediated by altered ion transport - comprises analysing sample for presence of mutation of potassium ion channel gene, ENAC, or in its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Primers AAV36488-V36500, and primers AAV40423-V40431 were used to amplify regions of the centrosomal protein 140 (CP140) gene from both normal and sclerotic individuals. To determine if patients have the disease, the obtained DNA from both types of individuals is subjected to standard PCR-RFLP. If any defects are identified at this loci, then the PCR fragments which contain the defective fragments are cloned and sequenced. The location of the defect can now be determined, characterisation of the
                                                                                                                                                                                                                            Example 1; Page 38;
                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1998-506740/43.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Epithelial sodium channel; ENAC; mutation; pathological condition; ion transport; water retention; blood pressure; metabolic acidosis; chronic respiratory disease; inflammation; human; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Exon 4 of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAV57641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 9 A; 4 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 31; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Screening for sclerotic disease in patients - by detection of anti-CP140 autoantibodies, altered CP140 mRNA, altered CP140 protein levels or an alteration in the CP140 gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-MAR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO9840516-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYYA ) UNIV YALB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2838 CAGGCAGCACAGATCAACATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  can occur,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     eapiene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      RP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              etandard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CAGGCAGTACAGATCAAGAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ENaC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     97US-0040171P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US004681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                thus allowing this gene to
                                                                                                                                                                                                                                                                                                                                                                                                                                                      ss,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            subunit amplifying forward primer B-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                         56pp; English.
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This invention describes a novel method for identifying an inhibitor of potentially useful for treatment of cancer, where the inhibitor is active con a gene vital for cell growth or viability, and where the gene is subject to loss of heterozygosity (LOH) in a cancer. The inhibitor is active considering the development of cancer in a patient having a precancerous condition, by administering to the patient a first allele concern in a cells of the precancerous condition, where the normal somatic cells of the patient are heterozygous for the first gene, the inhibitor concern is active on at least one but less than all allelic forms present in the population and trayets only one allelic form present in the commal somatic cells, and the first gene. The products and methods can be concern, atherosclerotic plaques, premalignant metaplastic or dysplastic concern, atherosclerotic plaques, premalignant metaplastic or dysplastic concern, benign tumours, endometriosis, polycystic kidney disease, and concern contends to the disease, and concern the contends can be concern, atherosclerotic plaques, premalignant metaplastic or dysplastic concern.
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Polymorphism; human; inhibitor; cancer; treatment; cell growth; LOH; cell viability; loss of heterozygosity; precancerous condition; ASI; allele specific inhibitor; somatic cell; diagnosis; prevention; atherosclerotic plaque; premalignant metaplastic lesion; endometriosis; dysplastic lesion; benign tumour; polycystic kidney disease; transplant; graft versus host disease; malignant cell removal; bone marrow; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Identifying target genes for allele-specific drugs - used for diagnosis, prevention and treatment of, e.g. cancers, atherosclerotic plaque, dysplastic lesions, endometriosis or graft versus host disease.
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                         immediately at the electrode surface. The microelectronic device is designed and fabricated to actively carry out and control reactions such as nucleic acid hybridizations, antibody/antigen reactions, sample preparation, diagnostics and biopolymer synthesis. The device can electronically control the transport and attachment of specific binding entities, such as nucleic acids and polypeptides, to specific microlocations. The device can subsequently control the transport and reaction of analytes or reactants at the addressed specific micro-locations. The device is able to concentrate analytes and reactants, remove nonspecifically bound molecules, provide stringency control for DNA hybridization reactions and improve the detection of analytes. The
                                                                                                                                                                                                                                                                                                                                                                The specification describes a self-addressable, self-assembling microelectronic device which is designed to actively carry out and control multi-step and multiplex molecular biological reactions in microscopic formats. A key aspect of this inventions is played by the ion permeable permeation layer which overlies the electrode. This permeation layer allows attachment of nucleic acids to permit immobilization but also separates the attached oligonucleotides and hybridized target DNA sequences from the highly reactive electrochemical environment generated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Microelectronic device; multi-step reaction; microscopic format; ion-permeable permeation layer; electrode; electrical control; transport; attachment; binding; DNA/RNA hybrid; probe; ss.
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RESULT 483
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Best Local S
Matches 17
                                                                                                                                                                                                                        for treating or preventing ischaemic diseases, especially ischaemic stroke. The invention also relates to methods of diagnosis, prognosis screening for genetic predisposition for diseases and disorders associated with increased levels of ANF. Pharmaceutical compositions, methods of screening for ANF alleles that are protective for stroke, methods of screening for ANF modulators useful for treatment and prevention of ischaemic stroke are also provided
                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                 Primers having the present sequence, and the sequence given in AAX25058, are used in the PCR amplification of the human atrial natriuretic factor (ANF) gene (see AAX25054) from a genomic or CDNA library. The invention provides methods for using mutant ANF (especially human mutant ANF in which Gly-1 is replaced by Ser, Thr or Tyr, see AAW98194-96), fragments, analogues, derivatives and homologues of mutant ANF proteins, the nucleic acids encoding these mutant ANF proteins, as well as modulators of ANF for the proteins of the second of the seco
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Homo sapiens.
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                                          5146 GGAACCATTTGCCTCTGGCTG 5166
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                                                                                                                                                                                  BP; 2 A; 5 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 13; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                               'Score 16.2;
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                                                                                Pred. No. 5.5
0; Mismatches
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                                                                                                                                                                                  U; 0 Other
                                                                                   3.8e+02;
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                                                                                                                                     Length 21;
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RESULT

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AAX26973/c
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RESULT 485
AAZ59350/C
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Matches 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   describes a protein comprising a mammaglobin antigen that is recognized by B and/or Tc cells specific for the natural, secreted and glycosylated form of mammaglobin polypeptide. This protein, or recombinant vectors that express it, are used in vaccines for treating mammaglobinexpressing cancers, specifically of the breast. Such cancers can also be treated using autologous tumor lymphocytes activated ex vivo with an mammaglobin antigen, then returned to the patient. Expression of mammaglobin is elevated in 27% of stage I primary breast cancers, so it represents a marker useful for diagnosis of this disease
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                                                                                                                                                                                                                   Single nucleotide polymorphism; SNP; STP2; phenol sulphotransferase; probe; genotyping; human; drug metabolism; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-APR-2000
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                                                                                                                                                                                                                                                                                                                                                   Human STP2 gene promoter polymorphism sequence 108.
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Local Similarity 85.7%;
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         Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 16.2; DB 1;
Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Indels
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AAZ44350/c
ID AAZ443
XX
AC AAZ443
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DT 04-APR
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Protei
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Antimi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequences AAZ59305-Z59352 are fragments of the human STP2 gene. The cragments are from the 8 exons, the promoter region, 3' and 5' untranslated regions of the STP2 gene. Each sequence contains a newly identified STP2 gene single nucleotide polymorphism (SNP). STP2 is a phenol sulphotransferase. Substrates for STP2 include minoxidil, acetaminophen, and paranitrophenol. Several of the nucleotide changes identified at the polymorphism sites, give rise to an amino acid changes may result in altered enzyme activity. The sequences can be used as probes for detecting STP2 polymorphisms. The polymorphic metabolism of STP2 substrates, potential drug-drug interactions and adverse side effects. They can also be used to detect diseases resulting from accidental or occupational exposure to toxins and to establish animal, cell or in vitro models for drug metabolism
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Best Local 9
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                                                           04-APR-1995;
                                                                                                                   04-APR-1995;
                                                                                                                                                                           07-DEC-1999
                                                                                                                                                                                                                                   US5998596-A.
                                                                                                                                                                                                                                                                                        Unidentified
                                                                                                                                                                                                                                                                                                                                             pathogenic microorganism; primer;
                                                                                                                                                                                                                                                                                                                                                                        Antimicrobial; cytostatic; immunosuppressive; protein prophylactic; therapy; treatment; cancer; autoimmune c
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Protein kinase inhibiting primer #12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAZ44350
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAZ44350 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 2 A; 9 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Page 17; 46pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel nucleic acid used for genotyping, e.g. to predict rate of drug metabolism.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUN-1998;
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                                                              95US-00416214.
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                                                                                                                   95US-00416214.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
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Pred. No. 6.
                                                                                                                                                                                                                                                                                                                                          cancer; autoimmune disease; ss.
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(USSH) US DEPT HEALTH & HUMAN SERVICES

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RESULT 487
AAZ75999
ID AAZ75999
XX AAZ759
XX Human
XX Human
KW Human
KW Genomi
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes novel purified aptameric oligonucleotides which have antimicrobial, cytostatic and immunosuppressive activity. The oligonucleotides are useful for binding to and preventing or inhibiting the biological function of a protein kinase or a target molecule and for detecting the presence or absence of a target molecule in biological samples. The oligonucleotides are also useful for prophylactic and therapeutic treatment of diseases such as cancer, autoimmune diseases and diseases caused by pathogenic microorganisms. This sequence represents a primer used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                             Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymotphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; dentification; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;
   AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention
                                                                                                                                                                                                                                               21-APR-1998;
23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                               diagnosis; ss.
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                                                                                                                                                                                                                                                                                           21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                       WO9954500-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human biallelic marker downstream amplification primer
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                                                                           Page 2438; 2745pp;
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98US-0109732P
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                                                                                                         genome.
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                                                                           English
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Pred.
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No. 6.
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                                                                                                                         density
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID
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                                                                                                                         disequilibrium
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RESULT 488
AAP96955/c
ID AAP969
XX AAP969
XX AAP969
XX Human;
XX Human;
XX Human;
XX Myocai
XX Myocai
XX Myocai
XX Myocai
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-SEP-1999;
26-JUL-2000;
16-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                               Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human gene single nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                (WHED )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphism; vascular disease; coronary artery
myocardial infarction; atherosclerosis; stroke;
pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; variant
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                                                                                                                                                                                                                                                                                                                                                                                WHITEHEAD INST BIOMEDICAL MILLENNIUM PHARM INC.
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; 2000US-0220947P.
; 2000US-0225724P.
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The present invention provides a method of diagnosing a vascular diseas in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin genes. The sequences at a number of polymorphic sites are also provided

vascular disease

provided

Example;

Page 163;

242pp;

English

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RESULT 489
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                                                          against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the
                                                                                                                                                                                                                                                                                              The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 101; Page 56; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vaccinating against tumors, infectious diseases, allergies and using immunostimulatory Py-rich and TG nucleic acids.
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27-SEP-1999;
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                                                                                                   moieties optionally extended by a polymerase. The branched chain compounds of the invention are used in nucleic acid synthesis reaction, primer extension reaction, reverse transcription reaction of RNA into DNA, nucleic acid hybridization experiment (for identifying sequence of nucleic acid), and nucleic acid amplification experiment (for analysing the expression pattern of genes). The compounds are also used in solid-phase enzymatic reactions. The present sequence was used in the course of the course of the present sequence was used in the course of the c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 10; 31pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (VBCG-) VBC GENOMICS GMBH.
                                                                            invention to produce branched chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAAAAAATACAAAAAAGAAAA 5413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hiller R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99EP-00125484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99EP-00125484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 0 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Huber M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "branch present"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "NH2-C6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "NH2-C6 attached"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ъ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G; 21 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 16.2; DB 1;
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                attached'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mueller
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 U; 0 Other;
                                                                                compounds
                                                                                                                                                                                                                                                                                                                                                                                  containing nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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                                                                                                                          œ.
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Query Match Best Local S Matches 18

Similarity

0.3%;

Score 16.2; DB 1 Pred. No. 6.8e+02

Length

21;

Indels

0

Gaps

0

Conservative

0

Mismatches

Sequence 21

BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;

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RESULT 491
AAF24290/c
ID AAF242
XX AAF242
XX AAF242
XX Comple
XX Comple
XX Comple
XX DNA ch
XX Uniden
XX U
RESULT 492
AAF86496
ID AAF8644
XX
AC AAF864
XX
DT 28-JUN
XX
DE PCR pr
XX
XX
Gene t
XX
CS
Synthe
XX
PO2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This involves the measurement of an electric current along the annealed strands compared to a standard. This is useful in the analysis of genet: polymorphisms and variation between genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Determining complementarity of nucleotide fragment for gene analysis, by comparing flow of electric current from or to electroconductive substrate through DNA fragment, with reference obtained from its complement.
  WO200123548-A1
                                                                                                          Gene therapy; RNase; DNA cleavage; enzyme; PCR primer; ss
                                                                                                                                                                                                                        28-JUN-2001
                                                                                                                                                                                                                                                                             AAF86496;
                                                                                                                                                                                                                                                                                                                               AAF86496 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention provides a method for analysing a nucleic acid strand to determine the degree of complementarity between two sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1; Page 12; 28pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Makino Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-JUN-2000; 2000EP-00112235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP1065278-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Complementary nucleic acid detection method related sequence #5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAF24290;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF24290 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-140003/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Complementary nucleic acid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA chip; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (FUJF ) FUJI PHOTO FILM CO LTD.
                                                                                                                                                                  primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5393 АЛАЛАЛАТАСЛАЛАЛАСАЛАЛ 5413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAAAAATACAAAAAAGAAAA 5413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAAAAAAAAAAAAAAAA 1
                                                                                                                                                                  M1G7173.rev used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Abe Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 1 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99JP-00159339
                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ogawa M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21
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                                                                                                                                                                                                                                                                                                                               BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 16.2; DB 1;
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Takagi M,
                                                                                                                                                                  construct mutant RNase P RNA construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           analysis; polymorphism; variation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Takenaka S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yamashita K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genetic
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RESULT 493
AAF86495/c
ID AAF864
XX AAF864
XX AAF864
XX PCR PI
XX Gene t
XX Synthe
XX W02001
XX W02001
XX W02001
XX Gene T
XX W02001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            片
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ś
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        that cleaves a DNA substrate with a Catalytic efficiency greater than wild-type RNase P RNA. The variant RNase P RNA is useful for cleaving a DNA substrate in a cell, which involves introducing the variant RNase P RNA and a guide sequence that hybridises to the DNA substrate of interest, into the cell, by introducing a vector encoding the variant RNase P RNA and the guide sequence as a single transcript, into the cell. Optionally RNase P protein subunit is also brought into contact with the DNA substrate of interest. The variant RNase P RNA is useful in diagnostic methods for detection, quantitation, or cataloging of DNA sequences, forensic methods, genome dissection methods, biostatistical
                                                                                                                                                                                                                    Gene therapy;
                                                                                                                                                                                                                                                                                                                                                      AAF86495 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        methods, and population genetics methods, cleaving genomic DNA at particular sequences, creating gene knockouts by gene cleavage, killing specific cells by specific cleavage of DNA, cleaving pathogen DNA in a host cell, and killing mutant cells by specific cleavage of mutant DNA in the cell. The present sequence is a PCR primer used in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Variant of RNAse P RNA molecule that specifically cleave DNA with catalytic efficiency greater than its wild-type, useful for DNA detection, quantitation or cataloging, biostatistical methods and
                                                  30-SEP-1999;
                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                       PCR primer M1G173.fwd used to construct mutant RNase P RNA construct.
                                                                                                                                                                                                                                                                                        28-JUN-2001
                                                                                                                                                                                                                                                                                                                         AAF86495;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 47; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-328183/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29-SEP-2000; 2000WO-US026864.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-APR-2001
                   (UYYA ) UNIV YALE
                                                                                 29-SEP-2000; 2000WO-US026864
                                                                                                                    05-APR-2001.
                                                                                                                                                   WO200123548-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nost cell, and necent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Dorit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYYA ) UNIV YALE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention relates to a variant RNase P RNA (see AAF86515),
                                                                                                                                                                                                                                                                                                                                                                                                                                                             4263 CTTCCACCTCTACCTGATCCC 4283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            CTTTCACCCTTACCTGATCCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cole KB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 3 A; 10 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                      RNase; DNA cleavage; enzyme;
                                                    99US-0156790P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0156790P
                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.3%;
                                                                                                                                                                                                                                                                                                                                                          21
                                                                                                                                                                                                                                                                                                                                                          ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 16.2;
Pred. No. 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                          21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
.8e+02;
                                                                                                                                                                                                                      PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
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RESULT 494
ABS78428/c
ID ABS784
XX ABS784
XX ABS784
XX Angiog
XX A
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA substrate in a cell, which involves introducing the variant RNase P RNA and a guide sequence that hybridises to the DNA substrate of interest, into the cell, by introducing a vector encoding the variant RNase P RNA and the guide sequence as a single transcript, into the cell. Optionally RNase P protein subunit is also brought into contact with the DNA substrate of interest. The variant RNase P RNA is useful in diagnostic methods for detection, quantitation, or cataloging of DNA sequences, forensic methods, genome dissection methods, biostatistical methods, and population genetics methods, cleaving genomic DNA at particular sequences, creating gene knockouts by gene cleavage, killing specific cells by specific cleavage of DNA, cleaving pathogen DNA in a host cell, and killing mutant cells by specific cleavage of mutant DNA in the cell. The present sequence is a PCR primer used in the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to a variant RNase P RNA (see AAF86515), that cleaves a DNA substrate with a catalytic efficiency greater than wild-type RNase P RNA. The variant RNase P RNA is useful for cleaving
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      catalytic efficiency detection, quantitati
                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABS78428 standard; DNA; 21 BP
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                                                                                                                                                         14-DEC-2000; 2000US-025534P.
                                                                                                                                                                                                                     14-DEC-2001; 2001WO-US048458
                                                                                                                                                                                                                                                                                                                                                                                                                                                              scleroderma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 47; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-328183/34
                                                                                                                                                                                                                                                                                                                                         WO200253141-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          angiofibroma; wound granulation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            torensic methods.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cell, and killing cell. The present sequence
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                                                                                                 COLEY PHARM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTTCCACCTCTACCTGATCCC 4283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cole
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inhibitory oligonucleotide #912.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypertrophic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ð,
                                                                                                    GROUP INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 scar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.2;
Pred. No. 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                intestinal adhesion; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3; Indels
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RESULT 495
ABL39404/c
ID ABL394
XX ABL394
XX ABL394
XX Immuno
XX Antibo
KW Angiog
XX Synthe
XX Synthe
YT W02001
PT 27-DEC
XX 22-JUN
XX 22-JUN
XX (IOWA
XX (IOWA
XX Winel
PT admin:
PT devel:
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PX GREET
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                  administering immunostimulatory nucleic acids that induce cell surface antigens and antibodies to a subject having
                                                    Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of
                                                                                                                                                                                                                                                                                                                                                                                     modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; phosphorothioate backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL39404
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Inhibiting angiogenesis in a subject, involves administering at least antiangiogenic nucleic acid molecule to the subject.
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                                                                                                      WPI; 2002-154611/20.
                                                                                                                                                                                                      22-JUN-2000; 2000US-0213346P
                                                                                                                                                                                                                                      22-JUN-2001; 2001WO-US020154.
                                                                                                                                                                                                                                                                                                          WO200197843-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Immunostimulatory nucleic acid SEQ ID NO: 840.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acid of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Page
                                                                                                                                                                                                                                                                         27-DEC-2001.
                                                                                                                                                                       (IOWA ) UNIV IOWA RES FOUND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAAAAAAAAAAAAAAAAA 1
                                                                                                                                      Hartmann
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       relates to inhibiting angiogenesis in
                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                          mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
                                                                                                                                                                                                                                                                                                                                         "phosphorothioate backbone"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 16.2; DB 1;
Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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                                        expression
or at risk (
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developing

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RESULT 496
AAD30438/c
ID AAD3044
AC AAD304
AC AAD304
AC AAD304
XX 21-MAY
XX 21-MAY
XX Human;
XW Human;
XW Prosta
XX POOR
CO OT-FEB
XX 27-JUL
XX 27-JUL
XX 27-JUL
XX 27-JUL
XX Chang
XX (UYRP
PF 27-JUL
XX Chang
XX (UYRP
PF 27-JUL
XX ASSESS
PT Subjec
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CC Cancer
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Best Local S
Matches 18
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                             The invention relates to a method for assessing the risk of prostate cancer in a human subject. The method involves determining the length of the contiguous CAG or CAA repeats in both AIBI (Amplified In Breast cancer 1) gene alleles or contiguous CAG, CAA or GGN repeats in the androgen receptor gene of the subject. The method is useful for assessing a subject's risk for acquiring or developing prostate cancer. The present sequence is a DNA encoding human androgen receptor (AR) polyglycine tract. This sequence is used in the molecular analysis and assessment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; AIB1; amplified in breast cancer 1; androgen receptor; AR; prostate cancer; polyglycine; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Assessing the risk of acquiring or developing prostate cancer in subject, comprises determining the length of the contiguous CAG, and/or GGN repeats in the AIB1 gene and/or androgen receptor gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-206195/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 309; 312pp; English.
                                                                                                                                                                                                                                                                                                                                                         Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-JUL-2000; 2000US-0221074P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYRP)
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                                                      crepeat within a coding sequence (expressed sequence tag, EST), which comprises detecting tandem repeats in a target coding sequence, scoring the repeats for polymorphic probability and generating a dataset correlating the repeats with polymorphic probability to identify a candidate polymorphic repeat. The computational methods (polymorphic marker prediction of ubiquitous simple sequences, POMPOUS, and Rep-X) are useful for identifying and detecting candidate polymorphic repeats in thuman genes, which can be used to understand, treat or eliminate genetic diseases, predispositions or adverse drug-treatment reactions. Examples of diseases linked to nucleotide repeats are Machado-Joseph, Haw River syndrome, Huntington's disease, fragile-X syndrome, Fredreich's ataxis, myotonic dystrophy, hyperandrogenaemia, spinal and bulbar attophy and spinocerebellar ataxia. The sequences presented in ABX79676-ABX80022 are the polymorphic repeats identified for a search of human ESTs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EST; expressed sequence tag; ss; polymorphic repeat; tandem rej polymorphic marker prediction of ubiquitous simple sequences; leep-X; human; genetic fleease; drug-treatment; Machado-Joseph; Haw River syndrome; Huntington's disease; fragile-X syndrome; Fredreich's ataxis; myotonic dystrophy; hyperandrogenaemia; spinal atrophy; bulbar atrophy; spinocerebellar ataxia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying a candidate polymorphic repeat within a coding sequence, understanding or treating genetic disease, comprises detecting tande repeats in a target coding sequence and scoring the repeats for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention discloses a method for identifying a candidate polymorphic
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polymorphic probability.
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Pred. No. 6.
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                                                                                                                                                         remote diagnostic server; processing digital sample signal at remotely located database to correlate digital signal with digital information and returning report of particular condition of animal. The method is useful for assessing a condition of a performance animal preferably human, dog or camel. The condition can be an athletic ability and a condition that enhances, hinders, impedes or does not change an expected ability of the performance animal; and also normal, pre-clinical, overt progress and/or stage of disease, undiagnosed of unclassified conditions, presence of drugs, response to exercise, response to vaccines, therapies, nutritional states and response to environmental conditions. Diseases assessed by the gastrittis, gastric ulcers, respiratory aliments, fractures, epistaxis, musculoskeletal damage or disorders and joint disease. The present sequence is a primer used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a method for assessing a condition of a performance animal. The method involves determining in sample abundance of expressed target nucleic acid; transmitting digital sample signal to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Assessing condition e.g. athletic ability, stage of disease, presence of drugs, response to exercise, response to vaccines, therapies, nutritional states, of performance animal involves analyzing nucleic acid expression.
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                                                                                                                             Sequence 21 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 46; 87pp; English.
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29-JUN-2001;
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2001US-00896941.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to stereoselective preparation of 2,3 dihydroxycarboxylic acids and derivatives and invovles hydrolysis of 2 epoxycarboxylic acids or derivatives in the presence of a polypeptide having epoxide hydrolase activity. The 2,3 dihydroxycarboxylic acids sused as chiral building blocks for more complex biologically active compounds. The reaction conditions are mild and the racemic epoxide starting materials are readily available. The present sequence represent appropriate for amplifying an epoxide hydrolase cDNA from R. araucari
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stereoselective preparation of 2,3-dihydroxycarboxylic acids, useful as chiral building blocks, comprises hydrolysis of 2,3-epoxycarboxylic acids in the presence of a polypeptide having epoxide hydrolase activity.
                                                                                                                             antiulcer;
psoriasis;
                                                                                                                                                                                                                                                                                         Immunostimulatory nucleic acid #881.
                                                                                                                                                                                                                                                                                                                                                                                25-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ACH03246;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ACH03246 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21
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Verdoes JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-SEP-2001; 2001EP-00120796
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-AUG-2003
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                                                                                      inflammatory bowel disease; ulcerative colitis;
                                                                                                                                                                                                                Immunostimulatory; antiinflammatory; dermatological; antipsoriatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-SEP-2001; 2001EP-00120796
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(KERJ ) FORSCHUNGSZENTRUM JUELICH GMBH.
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                                                                                                                             gene therapy; vaccine; non-allergic inflammatory disea eczema; allergic contact dermatitis; latex dermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       De Oliveira Villela Filho M, Liese A,
Weijers CAGM, Visser JH, Dreisbach
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 2 A; 8 C; 4 G; 7 T; 0 U; 0 Other;
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epoxide hydrolase; chiral building block; PCR primer; ss.
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Pred. No. 6.
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RESULT 501
ADB372
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ADB372
AC ADB372
AC ADB372
XX
O4-DEC
XX
OE Immuno
XX
OE; al
KW ds; al
KW hypo-x
XX
Synthe
XX
OS-PEB
XX
O2-PEB
XX
O3-PEB
XX
O3-PEB
XX
O3-PEB
XX
XX
DB (PETE/
PA (PETE/
PA (PETE/
PA (POUR)
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XX
DB BRATZ1
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              WPI; 2003-657977/62
                                          Bratzler RL,
                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                               ds; allergy; asthma; poly-G nucleic acid; aerosol formulation;
hypo-responsive subject; immunostimulatory.
                                                                                                                                                                                                                                                                                                                                                                                        ADB37209 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Treating non-allergic inflammatory diseases, such as psoriasis, ecz allergic contact dermatitis, latex dermatitis or inflammatory bowel
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                                                                                  (BRAT/)
(PETE/)
                                                                                                                          03-FEB-2000; 2000US-0179991P
                                                                                                                                                    02-FEB-2001; 2001US-00776479.
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                                                                                                                                                                                                                                                             nypo-responsive
                                                                                                                                                                                                                                                                                                     Immunostimulatory nucleic acid #823.
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                                                                                  BRATZLER R L.
PETERSEN D M.
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                                                                     FOURON
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                                          Petersen
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                                          Fouron
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 16.2;
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Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
Disclosure; Page 17; 221pp; English.
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FFXSXDDDDDDXS

Matches Query Match Best Local The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The methods and compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represents Sequence 21 an immunostimulatory nucleic acid of the invention. 18; Similarity BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other; Conservative 0.3%; <u>.</u> Pred. No. 6.8e+02; Mismatches Length Indels sequence represents 21; 0 Gaps 0

문 S 5393 АЛАЛАЛАТАСАЛАЛАДАЛАЛ 5413 AAAAAAAAAAAAAAAAA 1

21

ADJ92252; ADJ92252 standard; 06-MAY-2004 (first DNA; entry. 21

hair; keratin-associated protein; KAP; human; keratin; toiletry; therapeutic; hair growth promoter; hair disorder; PCR; primer; s Human hair keratin-associated-protein PCR primer P3 SEQ ID NO:111

Synthetic 22-MAY-2003 WO2003042387-A1

13-NOV-2001; 2001JP-00348050. 13-NOV-2002; 2002WO-JP011851. (UYKE-) UNIV KEIO.

JAPAN SOC PROMOTION SCI

DNAs encoding 39 Keratin-associated proteins localized on human chromosome 21, useful for screening binding and expression modias cosmetic and therapeutic agents for hair disorders. and expression modifiers and

WPI; 2003-493307/46.

Shibuya K,

Shimizu

z

(NIPR-)

Example 3; SEQ ID NO 111; 352pp; Japanese.

RESULT 502
ADJ92252
ID ADJ922
XX ADJ922
XX ADJ922
XX ADJ922
XX Homan
XX Hair;
KW therap
XX Homo (
XX Homo (
XX Homo (
XX I)3-NOV
XX I)3-NOV
XX II3-NOV
XX II3-NOV
XX II3-NOV
XX II3-NOV
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XX II3-NOV
XX IIA-NOV
XX The present invention describes DNAs encoding hair keratin-associated proteins (KAP) of human origin, which bind to hair keratin. Also described: (1) DNA encoding KAPs of mouse origin; (2) proteins encoded by the human and mouse DNA, and their partial peptides, and proteins derived from them by addition, deletion and/or substitution of one or more amino acid residues; (3) DNA hybridising to the DNA encoding KAP; (4) peptides SCCXPSCCXP (1), where X = Q, V, R or I, and YGAGYGSGY (II), where X = Y, L or F; (5) fusion proteins and peptides containing these proteins and peptides to the proteins and peptides; (3) recombinant proteins and peptides binding to these antibodies; (8) expression vectors containing the DNA encoding KAP; (9) host cells transformed by these vectors; (10) non-human animals which host cells transformed by these vectors; (10) knockout animals for KAP or which overexpress KAP; screening

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RESULT 503
ADKO1309/c
ID ADKO13
XX ADKO13
XX ADKO13
XX ADKO13
XX ADKO13
XX BAT DN
XX Rat DN
XX BS; hy
KW B1000d;
XX PEB
XX DE1020
XX PEB
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X
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Best Local Similarity
Matches 18; Conserv
This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture oligonucleotides are biotinylated and immobilised on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, metal, resin, gel, crystalline material and/or membrane, having semi-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      substances promoting or inhibiting the binding of KAP to hair keratin, (promoting or inhibiting the expression of KAP, comprising using KAP or their partial peptides or cells expressing them; (12) compounds identified by the screening method; and (13) tolletry and therapeutic compositions containing these compounds, or containing KAP or their partial peptides or (I) or (II), expression vectors for them, or host cells transformed by these vectors. KAP sequences can be used as hair growth promoters. The KAPs are useful as active ingredients in toiletry compositions (such as hair and beard growth improvers, hair colourants and hair conditioners) and in therapeutic compositions for hair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable and constant regions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ss; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rat DNA microarray
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 5 A; 10 C; 4 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-FEB-2002; 2002DE-01008794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-FEB-2002; 2002DE-01008794
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DEGUSSA BIOACTIVES GMBH.
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germ cell; food additive; food supplement.
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Pred. No. 6.8e+02.
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                                                                                                                                                                                                  conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. The method is used (i) for analysis of patterns, especially in mucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, and can detect very small differences in the nucleic acid gool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADX01281-ADX01344 represent
                                                                                Query Match
Best Local S
Matches 18
                                                                                                                                                                 Sequence 21
                                                                                                    Local Similarity
                                       5400 ТАСАЛАЛАЛАЛАЛАЛАТGAAA 5420
  21
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TACAAAAAAAAAAAAAAAA 1
                                                                                                                                                                 BP, 1 A, 0 C, 1 G, 19
                                                                                  Conservative
                                                                                                    0.3%;
85.7%;
                                                                                0
                                                                                                    Score 16.2; DB 1
Pred. No. 6.8e+02
                                                                                  Mismatches
                                                                                                                                                                     T; 0 U; 0 Other;
                                                                                                                            DB 1;
                                                                                                                          Length
                                                                                     Indels
                                                                                                                            21;
                                                                                     0
                                                                                     Gaps
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ADKO134/c
ID ADKO13
XX ADKO13
AC ADKO13
XX Set DN
XX Set DN
XX Set DN
XX Set DN
XX BS; hy
XX DE1020
PM D21020
PM D21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               capture oligonucleotide; cell; food additive; food
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d supplement.
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28-FEB-2002; 2002DE-01008794. 28-FEB-2002; 2002DE-01008794. 04-SEP-2003 (DEGS) DEGUSSA BIOACTIVES GMBH Ď Dieck HT, Hoppe Ξ

Sorting single-stranded nucleic acid, patterns and screening active agents, and constant regions. useful for analyzing expression uses capture agent with variable

WPI; 2003-714082/68.

Page 6; 8pp; German.

This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides have a 5'-invariable anchor region, the complement of which is present at least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region

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RESULT 505
ADKO1284/c
ID ADKO12
XX ADKO12
XX ADKO12
XX B5; hy
XX Ss; hy
XX Ss; hy
XX B100d;
XX DE1020
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XX DESEPE
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PT patte;
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PT patte;
CC readir
CC readir
CC compur:
CC compur:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cc comprises a sequence of 10-50, particularly 15-25, T residues. The cc capture oligonucleotides are biotinylated and immobilised on a surface by cinteraction with streptavidin. The matrix is of plassic, ceramic, glass, cc metal, resin, gel, crystalline material and/or membrane, having semi-cc conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single cc stranded nucleic acids to the surface is (quasi)covalent, supramolecular, physical, stimulated by an electrical field or through a molecular sieve. Cc The method is used (i) for analysis of patterns, especially in nucosal, cativity of pharmaceuticals and/or nutritional compounds, e.g. food cadditives or supplements, especially minerals, trace elements, organic mixtures. The method provides rapid, inexpensive and reproducible crepresentation of differences in pools of nucleic acid from cells. It callows imaging of the complete pattern of all nucleic acid fool. Since the comparison of nucleic acid pools, not individual capture probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                        Boekenkamp
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                                                                                                                                                                                                                                                                                                                                (DEGS ) DEGUSSA BIOACTIVES GMBH
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germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                        Dieck HT,
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Pred. No. 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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This invention describes a novel method for sorting single-stranded nucleic acids by isolation and hybridisation of nucleic acid pools, then reading out, where the nucleic acids are selectively bound using capture agents that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides

Sorting single-stranded nucleic acid, patterns and screening active agents,

useful for analyzing expression uses capture agent with variable

regions.

Page 4;

qq8

German

Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable

This invention nucleic acids !

ă

describes a novel method for a by isolation and hybridisation

sorting single-stranded of nucleic acid pools,

then

Example;

Page

6; 8pp; German.

constant

regions. screening 밁 Ś

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ADXO134/c
ID ADXO13
XX
AC ADXO13
AC ADXO13
XX
DT 06-MAY
XX
DB Rat DN
XX
Ss; hy
KW blood;
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Rattus
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CR
PD 04-SEP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               least once in each nucleic acid and a 3'-variable, discriminatory region that comprises all possible combinations of up to 10 nucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture agents are particularly locked nucleic acids (LNA) and the anchor region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ss; hybridisation;
blood; nerve; germ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADK01341 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                            (DEGS ) DEGUSSA BIOACTIVES GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA microarray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5390
                                                                                                                                                                                                                                                                                                2003-714082/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTAAAAAAATACAAAAAAGA 5410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 2 A; 0 C; 0 G; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   capture oligonucleotide; pattern; mucosal;
cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              capture oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 16.2;
Pred. No. 6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6.8e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     #61.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hair root;
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The state of the state of

Example; Page 5; 8pp; German.

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RESULT 507
ADK01329/c
ID ADK013
XX ADK013
XX ADK013
XX ADK013
XX AC ADK013
XX AC ADK013
XX BAL DN
XX BAL DN
XX BAL DN
XX BAL DS
XX DE1020
XX DE1020
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XX CDEGS
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Best Local Similarity
                                                                                               Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          04-SEP-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           06-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADK01329;
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                                                                                                                                                                                                                                                                                                                                            Boekenkamp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root;
blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADK01329 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                           (DEGS ) DEGUSSA BIOACTIVES GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5392 TAAAAAATACAAAAAAGAAA 5412
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                                                                                                                                                                                                                                                                                                                                            Dieck HT,
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85.7%;
                                                                                                                                                                                                                                                                                                                                            Hoppe H
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Pred. No. 6.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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RESULT 508
ADKO1336/c
ID ADKO13
XX ADKO13
XX ADKO13
XX ADKO13
XX O6-MAY
XX S8; hy
XX S8; hy
XX S8; hy
XX B1000i;
XX B2000i
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XX B2000i
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes a novel method for sorting single-stranded concleic acids by isolation and hybridisation of nucleic acid pools, then comparison that are (a) immobilised on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonuclectides chave a 5'-invariable anchor region, the complement of which is present at cleast once in each nucleic acid and a 3'-variable, discriminatory region comprises once in each nucleic acid and a 3'-variable, discriminatory region comprises all possible combinations of up to 10 nucleotides to allow that comprises all possible combinations of up to 10 nucleotides to allow binding of particularly locked nucleic acids (LNA) and the anchor region comprises a sequence of 10-50, particularly 15-25, T residues. The capture of interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface stranded nucleic acids to the surface is (quasi) covalent, supramolecular, compounds, particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi) covalent, supramolecular, compounds, especially minerals, especially in mucosal, that method is used (i) for analysis of patterns, especially in mucosal, cardiatives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and cardiatives. The method provides rapid, inexpensive and reproducible cards matrix miniaturisation of nucleic acid pool. Since the capture probes used in the method of the invention. ADX01281-ADX01344 represent contents on the particular acid in the method of the invention.
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Best Local S
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                                            WPI; 2003-714082/68
                                                                                                                Boekenkamp D,
                                                                                                                                                                                                                                                      28-FEB-2002; 2002DE-01008794.
                                                                                                                                                                                                                                                                                                                                 28-FEB-2002; 2002DE-01008794.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DE10208794-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ss; hybridisation; capture oligonucleotide; pattern; mucosal; hair root; blood; nerve; germ cell; food additive; food supplement.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rat DNA microarray capture oligonucleotide #56.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADK01336 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 2 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        capture probes used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5391 TTAAAAAATACAAAAAAGAA 5411
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                                                                                                                                                                                       DEGUSSA BIOACTIVES GMBH.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                   Dieck HT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.3%; Score 16.2;
85.7%; Pred. No. 6.
                                                                                                                Hoppe H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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RESULT 509
ADJ13120/c
ID ADJ131
XX ADJ131
AC ADJ131
XX Probe;
XX Probe;
XX Probe;
XX Lumoux
XX Homo 8
XX US2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Comprises a sequence of 10-50, particularly single entired that are solution with string out, where the nucleic acids are selectively bound using capture capture that are (a) immobilized on the surface of a solid matrix and (b) comprise variable and non-variable regions. The capture oligonucleotides capture as 5'-invariable anchor region, the complement of which is present at the capture oligonucleotides capture oligonucleotides capture oligonucleotides capture oligonucleotides to allow binding of particular sorts of single stranded nucleic acids. The capture oligonucleotides are biotinylated and immobilized on a surface by conducting properties are biotinylated and immobilized on a surface by interaction with streptavidin. The matrix is of plastic, ceramic, glass, conducting properties and especially in the form of a chip. Its surface is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, by single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, capture oligonucleotides of the surface is (quasi)covalent, supramolecular, conducting properties and especially in the form of a chip. Its surface capture is particularly a layer of (bio)molecular filaments and binding of single stranded nucleic acids to the surface is (quasi)covalent, supramolecular, chapture or gent root, blood, nerve or gent and conducting of single capture of biod, nerve or gent and capture and conducting of activity of pharmaceuticals and/or nutritional commonner. e.g. food
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               activity of pharmaceuticals and/or nutritional compounds, e.g. food additives or supplements, especially minerals, trace elements, organic acids (amino, carboxylic or fatty acid) or their derivatives, salts and mixtures. The method provides rapid, inexpensive and reproducible representation of differences in pools of nucleic acids from cells. It allows imaging of the complete pattern of all nucleic acid in a cell, are can detect very small differences in the nucleic acid pool. Since the method is based on comparison of nucleic acid pools, not individual genes, matrix miniaturisation is possible. ADX01281-ADX01344 represent
                                                                              27-JUN-2001; 2001US-0301370P
                                                                                                                                27-JUN-2002; 2002US-00184085
                                                                                                                                                                                                                                   US2003152950-A1
                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                           Human DNA probe used to immobilise CpG methylated DNA SeqID 247.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADJ13120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADJ13120 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sorting single-stranded nucleic acid, useful for analyzing expression patterns and screening active agents, uses capture agent with variable
                                                                                                                                                                                  14-AUG-2003
     (GARN/) GARNER H R. (MINN/) MINNA J D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5401 ACAAAAAAGAAAAATGAAAA 5421
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21
                                                                                                                                                                                                                                                                                                                                  ss; chemical modification; methylation; array; CpG island; suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probes used in the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 0 A; 0 C; 1 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   дB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 16.2; DB 1
Pred. No. 6.8e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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RESULT 510
ADM96310/cc
ID ADM963
XX ADM963
XX ADM963
XX ADM963
XX ADM963
XX ISOFOX
XX Human
XX Synthe
PN US2003
XX US2003
XX O9-MA3
XX O9-MA3
XX (SCHR/PA (CHEN/PA (SCHR/PA (S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             밁
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Garner HR, Minna JD,
   Enriching low abundance polynucleotides in a sample, useful for gene expression analysis, comprises exposing the sample to an enzymatically non-extendable nucleobase oligomer to block polymerase activity on high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human ATP5F1 gene, RT-PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             17-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADM96310;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; SEQ ID NO 247; 210pp; English.
                                                                                                                          WPI; 2003-901581/82.
                                                                                                                                                                                                                                                                                                                                                                     09-MAY-2002; 2002US-00144179
                                                                                                                                                                                                                                                                                                                                                                                                                           09-MAY-2002; 2002US-00144179.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US2003211483-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADM96310 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (LUEB/) LUBBKE K J. (BALO/) BALOG R P.
                                                                                                                                                                                                                                                                             (SCHR/) SCHROEDER B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      lsoform 1; ATP5F1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   human; H+ transporting; mitochondrial ATP synthase; subunit form 1; ATP5F1; reverse transcriptase; RT-PCR; primer.
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                                                                                                                                                                                                                                              SCHROTH G P.
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                                                                                                                                                                                  Chen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12 C; 0 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                         Schroth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       21
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Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                         GP
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abundance species

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RESULT 511
ADJ88057/c
ID ADJ880
XX ADJ880
XX ADJ880
XX O6-MAY
XX DE RT pri
XX Select
KW Select
XX W1den
XX Uniden
XX US2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC polymicalectide under conditions so that base pairing occurs, and conditions for polymerase extension. Preferably, CC the enzymatically non-extendable nucleobase oligomer does not have a CC ribose-containing oligomeric structure. It is a peptide nucleic acid CC (PNA) oligomer or is a modified nucleotide oligomer or internucleotide and condition or is a modified nucleotide oligomer in selected from 2'-cC modified and 3'-modified nucleotide oligomers in selected from 2'-cC modified nucleotide oligomers are selected from 2'-o-alkyl modified nucleotide oligomers are selected from 2'-o-alkyl modified nucleotide oligomers are selected from 2'-o-alkyl modified and 3'-alkyl modified nucleotide oligomers. The 2'-o-alkyl modified occurs are selected from 2'-o-alkyl modified nucleotide oligomers are 2'-o-alkyl modified analogue CC oligomers. The modified nucleotide oligomer or internucleotide analogue CC oligomers. The modified nucleotide oligomer is classes (MSP) oligomers, nucleotide oligomer or internucleotide analogue CC oligomers. Cl-C4 alkylphosphonate oligomer are methyl oligomers are methyl oligomers. The cl-C4 alkylphosphonate oligomers are methyl oligomers or internucleotides, beta-cC oligomers. The sample comprises more than one high abundance coligomer is chimeric. The sample comprises more than one high abundance comprises second strand cDNA synthesis. The sample is exposed to the nucleobase oligomer during the first and/or second strand cDNA synthesis. The sample is exposed to the nucleobase oligomer during the first and/or second strand cDNA synthesis. The sample is exposed to the comprises and amplification step, which is by comprises second strand cDNA synthesis. The sample is exposed to the comprises in a selective enrichment of low abundance in a comprise selective enrichment of low abundance of comprises in a selective enrichment of low abundance of comprises in a selectiv
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Best Local S
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    US2004014105-A1
                                                                                                                                                         RT primer used in the synthesis of an artificial gene transcript.
                                                                                                                                                                                                        06-MAY-2004
                                                                                                                                                                                                                                                                                              ADJ88057 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sample to an enzymatically non-extendable nucleobase oligomer having a nucleobase sequence complementary to a sequence within the high abundance
                                              Unidentified
                                                                                                               Selective enrichment;
                                                                                                                                                                                                                                                    ADJ88057;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       transporting,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 20; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      5393 ААЛЛАЛАТАСАЛАЛАДАДАЛА 5413
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    n 0.3%;
Similarity 85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mitochondrial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                               gene expression; RT; reverse transcriptase; primer;
                                                                                                                                                                                                                                                                                              21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            method of enriching a low abundance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATP synthase, subunit B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 16.2; DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred.
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Ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
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ADMO7216/c
ID ADMO72
XX
AC ADMO72
XX
AC ADMO72
XX
DE COntro
XX
DOUble
XW
DOuble
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CDNA 8
XW
CDNA 8
XW
CDNA 9
XX
CONTRO
XX
CON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enriching low abundance polynucleotide relative to a high abundance polynucleotide in a sample, for analyzing gene expression and creat cDNA libraries, comprises blocking polymerase activity on high abun
                                                                                                                                                (CHEN/)
(SCHR/)
(BRAN/)
(SCHR/)
                                                                                                                                                                                                                                                                           23-OCT-2002; 2002US-00278760
                                                                                                                                                                                                                                                                                                                         23-OCT-2002; 2002US-00278760
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                                                                                                                                                                                                                                                                                                                                                                                                                           US2004081962-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Double-stranded cDNA synthesis; cDNA first strand synthesis; cDNA second strand synthesis; RNA template; RNA amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Control primer used in cDNA first strand synthesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADM07216 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 0 A; 0 C; 0 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     artificial gene transcript.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to methods for the selective enrichment of low abundance polynucleotides. The invention is useful for analysing gene expression in a sample and creating cDNA libraries. The present sequence is reverse transcriptase (RT) primer used in the synthesis of an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      polynucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-MAY-2002; 2002US-00144179
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                                                   WPI; 2004-340131/31.
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                                                                                                                                                SCHROBDER B.
BRANDIS J.
SCHROTH G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SCHROTH G P.
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                                                                                                 Schroeder B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chen
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                                                                                                 Brandis J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Schroth
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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Pred. No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21 T;
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                                                                                                    Schroth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and creating high abundance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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Synthesizing double-stranded cDNA, by synthesizing a cDNA strand from

RNA

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RESULT 513
AAQ57215/c
ID AAQ572
XX AAQ572
XX AAQ572
XX AAQ572
XX AAQ572
XX BE Enzyma
XX Specif
KW Specif
KW Inhibi
KW Asthma
XX Synthe
XX WO9402
XX WO9402
XX WO9402
XX IN-JUI
PR 07-DEC
PR 07-DEC
PR 07-DEC
PR 07-DEC
PR 07-DEC
PR 19-JAN
XX Sullii
XX Sullii
XX Sullii
XX Sullii
XX Sullii
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CC mixture comprising reverse transcriptase, RNA template, and first strand CC primer complementary to template, removing the template, synthesising CC double-stranded cDNAs in a second reaction mixture comprising processive DNA polymerase, DNA ligase, first cDNA strand as template and random CC primers having a mixture of oligonucleotides having random DNA sequences. CC Also disclosed is a method for amplifying a population of RNA molecules comprising quantitative information about the generic product are useful in determining quantitative information about the generic profile of CC useful in differential gene expression assays for the invention is CC useful in differential gene expression assays for the analysis of CC diseased and normal tissue and for large-scale correlation studies on Sequences, mutations, variants or polymorphisms among samples. The method is efficient in synthesising improved cDNA molecules and effective in CC generating useful quantities of an amplified cRNA product that comprises a population of cRNA molecules in substantially the same relative molar cratio as the RNA or mRNA starting material. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                   07-DEC-1992;
07-DEC-1992;
19-JAN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    template, :
                                  Sullivan
                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                   inhibitor; inhibition; ribozyme; treatment;
asthma; inflammatory diseases; restenosis;
                                                                                                                                                                                                                                                                                                                                                                  Specific; cleavage; target RNA; protein; prophylaxis; expression; inhibitor; inhibition; ribozyme; treatment; prevention; psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and random
                                                                                                                                                     07-DEC-1992;
                                                                                                                                                                   17-JUL-1992;
                                                                                                                                                                                                     02-JUL-1993;
                                                                                                                                                                                                                                                                     WO9402595-A1
                                                                                                                                                                                                                                                                                                                                     hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                       Enzymatic
                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
26-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAQ57215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       stranded cDNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a method for synthesising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; SEQ ID NO 2; 19pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21
                                                                    RIBOZYME PHARM INC
                                    SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAAATACAAAAAGAAAA 5413
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21 BP; 0 A; 0 C; 0 G; 21 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                    RNA molecule stromelysin mRNA target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cDNA as template in the presence of processive DNA polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      removing the template and synthesizing double-stranded cDNAs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
                                    Draper KG
                                                                                                  92US-00916763.
92US-00987132.
92US-00989848.
92US-00989849.
93US-00008895.
                                                                                                                                                                                                                                                                                                                                     arthritis; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    by synthesising first cDNA strands in a first
                                                                                                                                                                                                     93WO-US006316.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.3%;
85.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    16.2; DB 1;
No. 6.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1;
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                                                                                                                                                                                                                                                                                                                                                                    psoriasis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
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RESULT 514
AAQ93472/c
ID AAQ934
XX AAQ934
XX AAQ934
XX AAQ934
XX Hammer
DT 06-DEC
XX Hammer
XX Hammer
XX Hammer
XX Hopati
XX H
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Best Local S
Matches 18
                                                             The sequences AAQ93462-Q93494 are examples of target cleavage sequences for a hammerhead ribozyme with sequence motif AAQ90453. A ribozyme, pref. hammerhead, halrpin, hepatitis delta virus, group 1 intron or RNAse P RNA motif can be used in a composition for the treatment of arthritis, cancer or angiogenesis. The ribozyme comprises between 5-45 bases complementary to the target mRNA. The ribozymes (see AAQ93830-51 for examples) were synthesised based on putative stromelysin mRNA target cleavage sequences (AAQ93496-Q93829). (Updated on 25-MAR-2003 to correct PN field.)
     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                       New enzymatic RNA molecules - which cleave mRNA of a gene encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hammerhead
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ93472;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ93472 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This is a stromelysin mRNA target sequence (nucleotide no. 455) of an enzymatic RNA molecule (ribozyme) which cleaves mRNA associated with the development or maintenance of osteoarthritis or other pathological conditions which are mediated by metalloproteinase activation. The concn. of the ribozyme necessary to effect a therapeutic treatment is lower than that of an antisense oligonucleotide and the specificity of action is higher. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Draper KG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9513380-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hammerhead ribozyme motif; arthritis; cancer; angiogenesis; hairpin;
hepatitis delta virus; group 1 intron; RNAse P RNA; stromelysin; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25-MAR-2003
06-DEC-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Enzymatic RNA inflammatory,
                                                                                                                                                                                                                                                                                                               Disclosure; Page 18; 70pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-NOV-1993;
                                                          (AAQ93496-Q93829). (Updated
                                                                                                                                                                                                                                                                                                                                                                                                                                                              1995-194099/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              104 CACCTCTTCTCAGCCTTGCAG 124
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                                                                                                                                                                                                                                                                                                                                                                      metalloproteinase, for treating arthritis, cancer or angiogenesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
        22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CACCTCTTCCCAGACTTTCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ribozyme target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 7 A; 2 C; 9
  BP; 7 A; 2 C; 9 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     molecules which cleave mRNA - used to treat or arthritic, stenotic or cardiovascular diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93US-00152487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94WO-US013129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence #11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 16.2;
Pred. No. 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>,</u>
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Length 22;

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RESULT 515
AAX63379/C
ID AAX6333
XX AAX633
XX Human
XX Human
XX Arthri
KW AAThri
HUMO 11-JUL
PR 23-DE
PR 
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The present invention describes a novel enzymatic nucleic acid (ENA) thaving a hammerhead motif (HM) comprising: (i) at least 5 ribose residues (; (ii) a 2'-C-allyl modification at position 4 of the ENA; (iii) at least ten 2'-O-methyl modifications; and (iv) a 3'-end modification. The ENA's can inhibit collagenase and stromelysin production in the synovial membrane of joints for the treatment or revention of arthritis, CC particularly osteoarthritis or rhemmatoid arthritis. The ENA's can also be used to treat antigen presenting cells of a donor to induce tolerance in a recipient to an alloantigen of a donor. They can also be used for cenhancing graft tolerance or for treating autoimnume disease, and for treating allergies and other inflammatory conditions. The ENA's can also be used in diagnosis. Ribozyme therapy impacts on the expression of stromelysin without introducing the non-specific effects upon gene cayression which accompany treatment with retinoids and dexamethasone.

The concentration of ribozyme required to affect a therapeutic treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
Matches 18; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-DEC-1994;
23-DEC-1994;
23-DEC-1994;
17-PEB-1995;
20-APR-1995;
02-MAY-1995;
04-MAY-1995;
07-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beigelman
Mcswiggen
Karpeisky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Enzymatic nucleic acid molecules having a hammer-head motif -
the treatment of arthritis, induction of graft tolerance or to
auto-immune diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9618736-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Arthritic condition; graft tolerance; immune response; target; cl hammerhead ribozyme; hairpin ribozyme; human; rabbit; mouse; coll stromelysin; synovial membrane; joint; arthritis; osteoarthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX63379 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 139; 307pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-300653/30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rheumatoid arthritis; autoimmune disease; allergy; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human stromelysin hammerhead target
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIBO-) RIBOZYME PHARM INC.
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Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Stinchcomb DT,
Gustofson J, U
Thompson JD, M
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94US-00363253
94US-00363264
95US-00390850
95US-00426124
95US-0043674
95US-00434509
95US-0000951P
95US-0000974P
95US-00512861
95US-00541365
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Usman N, Wincott F, Modak A, Burgin A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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0; Mismatches 3
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Matulic-Adamic
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    used for
treatment

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at least
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RESULT 51

ABK64865

ID ABK6

AC ABK6

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                                                          Query Match
Best Local S
Matches 18
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                   The invention relates to a new antisense compound (1) up to 50 nucleobases in length and which specifically hybridises with a nucleic acid encoding breast tumour amplified kinase (BTAK) and inhibits the expression of BTAK. (1) is useful for inhibiting the expression of BTAK in human cells or tissues and for treating an animal, particularly a human suspected of having or being prone to a disease or condition associated with expression of BTAK. (1) is useful for diagnostics, therapeutics, prophylaxis and as research reagent and kits. (1) is useful prophylactically to prevent or delay infection, inflammation or tumour formation. ABK64961-ABK6496 represent human BTAK coding sequences, antisense oligonucleotides and related primers of the invention
                                                                                                                                                                       Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compounds targeted to nucleic acids encoding breast tumor amplified kinase, useful for modulating gene expression and treating diseases associated with expression of breast tumor amplified kinase in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           is lower than that required of antisense molecules, and is highly specific. The present sequence is used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cowsert LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antiinflammatory; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; BTAK; breast tumour amplified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human breast tumour amplified kinase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-JUN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               516
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     162 GGGAAGAATCTGAGGAACACA 182
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                                                                                        Similarity
                                                                                                                                                                       BP; 9 A; 2 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Col 42; 35pp; English
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                                                                                        0.3%;
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                                                             0
                                                                                     Score 16.2; DB 1
Pred. No. 6.9e+02
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Pred. No. 6.9e+02
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                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BTAK) reverse PCR primer.
                                                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic;
                                                                                                               Length 22;
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                                                             Gaps
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RESULT 518
ADD694
ID ADD694
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AC ADD694
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DT 15-UAN
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DT 5' anc
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Inter
KW inter
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Synthe
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AAL60163
ID AAL601
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 9 A; 5 C; 7 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                       The invention relates to a novel polypeptide, termed Novel Gene Expresse in Prostate (NGEP), which is detected in the cells of prostate (NGEP) antibodies are useful in detection assays, in the production of immunoconjugates, such as immunotoxins, which can be used to target prostate cancer. Polypucleotide encoding NGEP, or an NGEP polypeptide can be used for treating prostate cancer and generate an immune response against prostate cancer cells. The present sequence is a PCR primer for amplification of human prostate-specific EST (expressed sequence tag)
    WO2003085133-A2
                                                  inter-simple sequence repeat; ISSR; SSR, PCR;
animal; Basmati rice; ss.
                                                                                          5' anchored
                                                                                                                                             ADD69448;
                                                                                                                                                                     ADD69448 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 2; Page 44; 74pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New isolated Novel Gene Expressed in Prostate polypeptide and encoding nucleic acid molecule, useful for diagnosing and treating prostate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2003-449573/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pastan
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expressed s
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                                                                                                                    15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-MAY-2003
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                                                                                                                                                                                                                                    AGCTAGACAGGCAGCAGGACA 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             e Expressed in Prostate; NGEP; prostate cancer; human; EST;
sequence tag; PCR primer; ss.
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                                                                                          (ISSR)-PCR
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                                                                                                                  (first entry)
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                                                                                                                                                                     BP.
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                                                                                                                                                                                                                                                                                               Score 16.2; DB 1;
Pred. No. 6.9e+02
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                                                                                         SEQ
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                                                                                          ID 6.
                                                                                                                                                                                                                                                                                                             DB 1;
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                                                                primer;
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                                                                                                                                                                                                                                                                                                            Length 22;
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                                                              genotyping;
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RESULT 519
ACHO0645
ID ACHO06
XX ACHO06
XX ACHO06
XX INVERT
DT 12-FEE
XX INVERT
CS Mammal
CS MAM
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 22 BP; 10 A; 0 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a novel set of inter-simple sequence repeats (ISSR)-PCR primers for genotyping eukaryotes. The primers of the invention may be useful for genotyping diverse genomes of plant and animal systems, in particular for distinguishing Basmati rice varieties from non-Basmati rice varieties and traditional Basmati rice varieties from evolved Basmati rice varieties. The current sequence is that of the
                                                                              Disclosure; Page 23; 63pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ACH00645 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New set of inter-simple sequence repeats (ISSR)-PCR primers i
genotyping eukaryotes, useful for genotyping diverse genomes
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                                                                                                                            Determining predisposition breeding animals comprises
                                                                                                                                                                                                                                                               Hardge T,
                                                                                                                                                                                                                                                                                                                                                                      05-FEB-2002; 2002EP-00002632
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inverted nipple; microsatellite; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mammalian inverted nipple associated microsatellite PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        animal systems.
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                                                                                                                                                                                                                                                                                                                     (FOER-) FOERDERVEREIN BIOTECHNOLOGIEFORSCHUNG
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                                                                                                                                                                                                                                                               Schellander K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ISSR)-PCR primer of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22
                                                                                                                            to inverted nipples useful e.g. for selecting detecting specific microsatellite markers.
                                                                                                                                                                                                                                                                  Wimmers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 16.2;
Pred. No. 6.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer
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The present invention the predisposition of

relates to appearance

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use of a nucleic acid to determine inheritance of inverted nipples,

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RESULT 520
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The invention relates to a method of designing primers for simultaneous camplification of target DNA fragments in a single multiplex polymerase can amplification of target DNA fragments in a single multiplex polymerase can all primes as first primer and a second primer. The method comprises: (a) aligning a first primer and a second primer, and (b) contain four or more bases that are perfectly matching to the 3' end does not contain four or more bases that are perfectly matching to the 3' end cos not contain seven or more bases that are perfectly matching careful careful one mismatch to the 3' end sequence of the first primer or the second primer, the first primer at its 3' end does not contain six or careful primer or the second primer at its 3' end does not contain six or the second primer or the second primer at its 3' end does not contain six or the second primer at the siret primer at its 3' end consistant to a sequence anywhere of the first primer or the second cone mismatch to a sequence bases that are perfectly matching except one mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch to a sequence anywhere of the first primer or the second cone mismatch or the second cone mismatch to a sequence anywhere of the first primer of the second cone mismatch or the second cone mismatch or t
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Single multiplex PCR primer #1748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 41; 120pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            multiplex DNA sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Designing primers for simultaneous amplification of target DNA fragments in a single multiplex polymerase chain reaction, for high throughput
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-340914/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (UYNE-) UNIV NEW JERSEY MEDICINE & DENTISTRY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3419 AGATGAGGCAGGAACTGAGGG 3439
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        le multiplex polymerase chain reaction, DNA sequence amplification, comprises
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85.7%;
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Pred. No. 6.9e+02;
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08-APR-1991;
17-APR-1991;
17-APR-1991;
17-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   chain reaction. It is also useful in the identification of multiple genes related to multifactorial diseases, the genome-scale detection of genetic alterations, the studies in pharmacogenetic reactions, the genotyping genetic polymorphisms in a large population, the gene expression profiling in various samples and high throughput genotyping technologies. This sequence corresponds to an example of a primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human interleukin-6 gene; herpes simplex; AIDS; modified; HIV; RSV; HPV; malignancy; hepatitis; inflammation; ss.
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                                                                     WPI; 1992-217083/26.
                                                                                                                                                                                                                                                                                                                                                      23-NOV-1990;
                                                                                                                                                                                                                                                                                                                                                                                                    25-NOV-1991;
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07-DEC-1992
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                                                                                                                                                                 (GILE-) GILEAD
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ilarity 85.7%;
Conservative
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91US-00643382.
91US-00683420.
91US-00686544.
91US-00686546.
91US-00686547.
                                                                                                                                                                 SCI INC.
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/mod_ba
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/label= inverted_polarity_region
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                                                                                                                   Matteucci
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Pred. No. 6.
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                                                                                                                      Milligan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 22;
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New oligomers contg. modified bases - which form a triplex with G-C doublet in a DNA duplex, for treating and diagnosing HIV, hepatitis.

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RESULT 522
AAX57200/c
ID AAX57200;
XX
AX57200;
XX
AXC AAX57200;
XX
DT 26-JUL-1999 (first XX)
CCT-1999 (first XX)
DT 26-JUL-1999 (first XX)
DT 26-JUL
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Matches
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      This sequence describes novel methods where a genetic selectable construct operatively linked to a porcine promoter is introduced source of cells that contain porcine stem cells under conditions allows for differential expression of the marker and hence the se
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oct-4; porcine; selectable marker construct; promoter; stem cell; differential expression; selective isolation; pluripotent; propagation; cell surface membrane protein; rejection; xenotransplantation; transgenic pig; organ; transplant; programmed cell death; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Porcine Oct-4 PCR primer 6-T7 OL.4 5'.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 23
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                                                                                                                                                                                                                                                                                                                                                                                                                      (BIOT-) BIOTRANSPLANT INC.
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                                                                                                                                                               Page 28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry
                                                                                                                                                                                                                             cells comprising a marker gene under
                                                                                                                                                                                                                                                                                                                                                        Akiyoshi DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         71;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
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Pred. No. 6.9e+02;
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RESULT 523
AXX81611
ID AXX816
XX AXX816
XX AXX816
XX AXX816
XX Exythx
XX Exythx
XX Synthe
OS Erythx
XX Synthx
XX FR2771
XX FR2771
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XX INGUSE
PR 03-DE(
XX O3-DE(
XX INGUSE)
PR (ASSI-
XX INGUSE)
PR (ASSI-
XX INGUSE)
PR WPI; INGUSE
XX INGUSE
PR WPI; INGUS
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Query Match
Best Local Similarity
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Best Local
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                                                                                                                                    AAX81588-X81630 represent PCR primers used to amplify erythrovirus V9 polynucleotide sequences. Probes and primers derived from erythrovirus polynucleotide sequences (AAX81580) can be used for differential diagnosis of erythrovirus (parvovirus) infections by a combination of amplification and hybridiaation assay. The probes can also be used to assess susceptibility to erythrovirus infection and for erythrovirus screening and typing. The antibodies can be used in immunoassays for diagnosis of erythrovirus V9 infections
                                                                                    Sequence 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23 BP; 8 A; 8 C; 3 G; 3 T; 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Erythrovirus V9 and its nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-349543/30.
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                                                                                                                                                                                                                                                                                                                                                                                                            Page 27; 80pp; French.
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                                                                                       BP; 16
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                                                                                       A; 1 C; 1 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Auguste
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Pred. No. 6.
Score 16.2; DB 1;
Pred. No. 6.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acid sequences -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U; 1 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleotide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
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                           Length
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RESULT 524
AAI70049/c
ID AAI700
XX AAI700
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XX Synthe
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XX EP113;
XX I1-MAI
PF 06-MAI
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PT New t]
PT motif
RESULT 525
ABL91058
ID ABL910
XX
AC ABL910
XX
DT 27-WAY
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DE Homini
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Best Local S
Matches 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to thermostable mutant B-type DNA polymerases, which have a Y-GG/A amino acid motif between the N-terminal 3'-5' exonuclease domain and the C-terminal polymerase domain, where the tyrosine of this motif is mutated. The mutant B-type DNA polymerase is useful for synthesising nucleic acids and for PCR. To illustrate the invention, Thermococcus aggregans (Tag) DNA polymerase was used (AAI70053). The present sequence is a PCR primer, which was used in an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New thermostable mutant B-type DNA polymerase with a Y-GG/A amino acid motif between the N-terminal 3'-5' exonuclease domain and the C-terminal polymerase domain in the wild type polymerase, useful for synthesizing
  Hominidae; low density lipoprotein receptor; LDL receptor; LDL-R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 23 BP; 4 A; 8 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 5; Page 9; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-MAR-2001; 2001EP-00104583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-SEP-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAI70049;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAI70049 standard;
                                                                                                        27-MAY-2002
                                                                                                                                                            ABL91058
                                                                                                                                                                                                            ABL91058 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-618367/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11-MAR-2000; 2000EP-00105155
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer lambda 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                13-FEB-2002
                                                Hominidae LDL receptor related DNA sequence #102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (HOFF ) ROCHE DIAGNOSTICS GMBH
                                                                                                                                                                                                                                                                                                                                                                        4406 AGAAAGATGAGACTCTGGTGT 4426
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5400 TACAAAAAGAAAAATGAAA
                                                                                                                                                                                                                                                                                                                                        21
                                                                                                                                                                                                                                                                                                                                                                                                                                                 18;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                           ATAAAGATGAGACGCTGGAGT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Frey B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 16.2;
Pred. No. 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 526
AAPOS471/c
ID AAPO54
XX AAPO54
XX Hammer
XX Hammer
XX Hammer
XX Homo B
XX H
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes a method for detecting lipid metabolism errors in patients using as indicators a set of 65 specific low density lipoprotein (LDL) receptor gene mutations. The method can be used in the diagnosis of an inherited predisposition to the development of diseases associated with hyperlipaemia, such as arteriosclerosis and ischaemic heart disease. ABL91141 encodes the LDL receptor given in ABB90525. ABL91142 to ABL91183 represent PCR primers used in the amplification of the receptor gene. ABL90990 to ABL91140 and ABB9045 to ABB90524
                                                                                                               12-APR-1999;
                                                                                                                                                                                                                           19-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                 Ribozyme; erythropoietin; granulocyte colony stimulating factor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hammerhead ribozyme substrate #2690.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF05471
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      represents sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Set of specific low density lipoprotein receptor gene mutations for diagnosis of familial lipid metabolism errors including hyperlipemia.
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Iwasaki T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-JUL-2000; 2000JP-00218039
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                                                                                                                                                                     11-APR-2000; 2000WO-US009721
                                                                                                                                                                                                                                                                              WO200061729-A2
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                                                        (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BMLB-) BML INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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78.3%;
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Pred. No. 6.
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Blatt L,

Zwick M,

Pavco P,

Mcswiggen

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RESULT 527
AAF05469/c
ID AAF054
XX AAF054
XX AAF054
XX Hammer
XX Ribozy
XW Ribozy
XW Ribozy
XW Interf
XX Homo 8
XX Tl-APR
XX HI-APR
XX HI-APR
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CC molecu
CC Inhibi
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Best Local Similarity
Matches 16; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Enzymatic useful for
                                The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, EAR3/COUP-TF-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                            Enzymatic and antisense nucleic acid inhibition of repressor genes useful for producing e.g. granulocyte colony stimulating factor printerferon alpha and erythropoietin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to enzymatic and antisense nucleic acid molecules that act as inhibitors of the expression of repressor genes encoding the TR2 Orphan receptor, ER3/COUP-TP-1, the GATA transcription factor gene, IRF-2 and/or the CAATT Displacement Protein (CDP). Inhibition of the repressors removes prevents inhibition (and consequently increases expression of) genes involved in the production of erythropoietin, granulocyte colony stimulating factor protein and
                                                                                                                                                                                                                                                                                                                   Blatt L,
                                                                                                                                                                                                                                                                                                                                                                                        12-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                           11-APR-2000; 2000WO-US009721
                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ribozyme; erythropoietin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hammerhead
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAF05469;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF05469 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 17 BP; 2 A; 1 C; 1 G; 13 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 18; Page 117; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             useful for producing e.g. granulocyt interferon alpha and erythropoietin.
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                                                                                                                                                                                                                                                                                                                                                      (RIBO-)
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                                                                                                                                                                             18;
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                                                                                                                                                                                                                                                                                                                                                    RIBOZYME PHARM
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                                                                                                                                                                          Page 117; 164pp; English.
                                                                                                                                                                                                                                                                                                                   Zwick M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ribozyme
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense nucleic acid inhibition of repressor genes, oducing e.g. granulocyte colony stimulating factor pro
                                                                                                                                                                                                                                                                                                                                                                                        99US-0129390P
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                                                                                                                                                                                                                                                                                                                   Pavco P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  substrate
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Pred. No.
                                                                                                                                                                                                                                                                                                                  Mcswiggen
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7.1e+02;
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RESULT 52
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Best Local S
Matches 16
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                 modified_base
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cancer; cardiant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antisense therapy; infection; cardiovascular disorder; immune reaction; gene therapy; virucide; cytostatic; antibacterial; antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ON-21 oligonucleotide used
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                                                                                                                                                                                                                                                                                                                                                                                                                  Key
                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified.
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68/c
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ilarity 100.0%;
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                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "5-(1-propynyl)-2'-deoxyuridine;
given as N in the sequence shown as SEQ
sequence listing"
9
                        /note= "5-(1-propynyl)-2'-deoxyuridine; given as N in the sequence shown as SEQ sequence listing"
                                                                                    N in the sequence
                                                                                                                                   /mod_base= OTHER
/note= "5-(1-propynyl)-2'-deoxyuridine;
given as N in the sequence shown as SEQ
sequence listing"
                                                                                                                                                                                  listing"
11. .16
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/note= "5-(1-propynyl)-2'-deoxyuridine; This base
given as N in the sequence shown as SEQ ID NO: 15
                                                                                 /mod_base= OTHER
/mode= "5-methyl-2',-deoxycytidine; This base
/note= "5-methyl-2',-deoxycytidine; This base
/ n in the sequence shown as SEQ ID NO: 15 in t
                                                                                                                                                                                                                     /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                            sequence
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                                                     /*tag=
/mod_bas
                                                                                listing
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/note= "5-methyl-2'-deoxycytidine;
/note= "5-methyl-2'-deoxycytidine;
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mod_base=
         *tag=
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                                                     _base= OTHER
                                                                                                                                                                                                                       base= OTHER
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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7.1e+02;
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NO: 15 in t
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ID NO: 15
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0: 15
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sequence
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sequence
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sequence
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RESULT 529
ABZ10676/c
ID ABZ106
XX ABZ106
XC ABZ106
XC ABZ106
XC ABZ106
XC Haemat
XC Haemat
XC Human;
KW Gytosi
XX Homo g
OS Synthe
XX Synthe
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Best Local Similarity
Matches 16; Conserv
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25-AUG-1992;
23-OCT-1992;
25-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to novel oligomers which have enhanced ability with respect to forming duplexes or triplexes. The oligomers comprise at least three 3'-5' linked nucleosides or their salts. At least one internucleoside linkage is not a phosphodiester linkage and at least one nucleoside comprises a base. Sequences of the invention are useful for binding to a DNA duplex target sequence via either CT or GT triplex helix binding motif and in antisense therapies. They are also used for treating diseases caused by viruses and for diagnostic applications to detect viral infections, bacterial infections and diseases such as cancers. The oligomers are also used as primers, in the treatment of pathological conditions associated with inflammatory conditions, cardiovascular disorders, immune reactions and bacterial infections and to modulating target gene expression. They are also useful in gene therapy. The present sequence is an oligonucleotide used in the
                                                              Human; haematopoietic cell proliferation disorder; cytostatic; gene therapy; lymphocytic leukaemia; acute myelogenous leukaemia; cytosine methylation state; probe; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligomers useful for binding to DNA duplex target sequence and for treating e.g. diseases caused by viruses and inflammatory conditions comprise at least three 3'-5' linked nucleosides.
                 Homo sapiens
Synthetic.
                                                                                                                                                                  16-JAN-2003
                                                                                                                                                                                                   ABZ10676;
                                                                                                                                                                                                                                ABZ10676 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 17 BP; 2 A; 3 C; 0 G; 0 T; 12 U; 0 Other;
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                                                                                                                                 Haematopoietic cell proliferation disorder related oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Froehler B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PHARM INC.
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92US-00935444.
92US-00965941.
92US-00976103.
94US-00338352.
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                                                                                                                                                                                                                                                                                                                                                                                                  0.3%;
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Pred. No.
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7.1e+02;
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ARESULT 530
ADB65475/C
ID ADB654
XX ADB654
XX ADB654
XX ADB654
XX B9-JAN
XX Human
XX Human
XX Theuman
XX Theuma
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XX Income 88; pr
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dinucleotides within the target nucleic acid. ABZ09861 to ABZ1118
CC represent specifically claimed nucleotide sequences from the present
CC invention. Oligonucleotides from the present invention can be used: for
CC differentiating between healthy haematopoietic cells and proliferative
CC disorder haematopoietic cells; for differentiating between acute
CC disorder haematopoietic cells; for differentiating between acute
CC pymphocytic leukaemia and acute myelogenous leukaemia; as probes for
CC determining the cytosine methylation state and/or single nucleotide
CC polymorphisms (SNPs) of haematopoietic cell proliferation disorder
CC related sequences and their complements; and as primers for the
CC amplification of haematopoietic cell proliferation disorder related DNA
CC sequences. The nucleotide sequences from the present invention can also
CC be used for detecting a predisposition to, differentiation between
CC used for detecting a predisposition to, differentiation between
CC highly specific cell proliferative disorders. The present method enables a
CC highly specific classification of haematopoietic cell proliferative
CC disorders allowing for improved and informed treatment of patients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local (
                                                                                                                                                                                                                    Human WNT3 reverse PCR primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            differentiating between haematopoietic cell proliferative disorders associated with at least I gene and/or their regulatory regions in a subject. The method comprises contacting a target nucleic acid in a biological sample obtained from the subject with at least I reagent,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WO200277272-A2
                                                                                                                     ss; primer; human; PCR; WNT; chronic rheumatoid arthritis; WNT10B;
rheumatoid arthritis; osteoarthritis.
                                                                                                                                                                                                                                                                                      29-JAN-2004
                                                                                                                                                                                                                                                                                                                                                    ADE65475
                                                                                                                                                                                                                                                                                                                                                                                                            ADE65475 standard;
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Mueller V,
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WO2003093508-A1 Homo sapiens

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                                                                     Use of and/or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a novel method for diagnosis of chronic rheumatoid arthritis in which synovial fluid, joint tissue or peripheral blood is analysed to detect greater than normal expression of WNT10B. The method is useful for simple diagnosis of rheumatoid arthritis and its discrimination from osteoarthritis. The present sequence represents a PCI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-inflammatory; neuropeptide Y Y1 receptor; NPY Y1; antagonist; cutaneous; internal inflammation; neurogenic; acute; chronic; drug target; 129SVXBalb/c hybrid mouse; pain transduction; PCR pri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unidentified.
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treating inflammatory c
g target for screening a
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  Page 11;
                                                                                                                                                                                       Naveilhan
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Pred. No.
                                                antiinflammatory
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Mismatches
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                                                                     for preventing inflammation
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The present polynucleotide sequence represents the PCR primer 3 for use in analysing Y1 mutant 1298VXBalb/c hybrid mice. The mutant hybrid mice were created by the injection of a Y1 targetting construct into homologous recombinant embryonic stem cell clones. The invention relates to the use of, or method of using, a selective neuropeptide Y Y1 (NPY Y1) receptor agonist for preparation of a drug for preventing and/or treating pain conditions. In a further aspect the invention relates to use of the NPY Y1 receptor as a drug target in screening procedures to find agonists of said receptor, more precisely to find anti-nociceptive compounds which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          neuropeptide Y Y1 receptor agonist; anti-nociceptive; pain alleviation; cutaneous pain; visceral pain; chemical pain; thermal pain; diffuse pain; mechanical pain; local pain; chronic pain; persistent pain; dorsal horn;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         primer 3 for analysis of Y1 mutant 129SVXBalb/c hybrid mice.
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                                                                                                                                                                                                                                                                                                                                                                         lective neuropeptide Y1 for treating pain e.g.
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                                                                                                                                                                                                                                                                                           Page 11; 30pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             neurogenic inflammation or acute and chronic The method is further useful as drug target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     find anti-inflammatory compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     spinal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              De Araujo Lucas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 16; DB; Pred. No. 7.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ΑB
                                                                                                                                                                                                                                                                                                                                                                                     receptor agonist diffuse or local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U; O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Le
7.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ရှ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hassani
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                  in the preparation of or chronic/persistent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21
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RESULT 533
AAQ51663/c
ID AAQ516
XX AAQ516
XX AAQ516
XX AAQ516
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XX ADV pr
XX ADV; A
KW ADV; A
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PA (ZENK-
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XX WPI; 1
XX Detect
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XX Detect
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RESULT 534
AAT30413
ID AAT304
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                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 17
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                         The detection method for Aujezsky's disease virus uses a primer pair selected from the ADV DNA-specific region, with PCR designed to suit the amplification of DNA, thereby permitting specific and highly sensitive detection of ADV. Claimed primers are given in AAQS1659-72; additional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               directly or indirectly affect the NPY Y1 receptor in a selective way for treatment of the various pain conditions. The NPY Y1 receptor agonist may be administered to alleviate cutaneous, visceral, chemical, thermal and mechanical pain conditions, also to alleviate diffuse or local pain conditions, and to alleviate chronic/persistent pain conditions. Neuropeptide Y is believed to exert anti-nociceptive actions by inhibiting the release of substance P and other pain neurotransmitters in the dorsal horn of the spinal cord
                                                                                                                                                                                                                                                                                                                                                Sequence 19 BP; 2 A; 6 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1993-373607/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (NISS )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           JP05276998-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADV; Aujeszky's disease virus; primer; PCR; amplification; polymerase chain reaction; detection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADV primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ51663;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAQ51663 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26-0CT-1993
                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tection of Aujeszky's disease virus primer for selective detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1306 AGCCTCTGTTCCACAT 1321
                                                                                                                                                                                      3251 GCTGCCAGGACCTGGCCTC 3269
                                                                                                                                                                                                                                           17;
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                                                                                                                                                                                                                                                                                                                                                                                                   are given in AAQ51673-74
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 1; 9pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGCCTCTGTTCCACAT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (III)a.
                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92JP-00079881
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NOGYO KYODO KUMIAI REN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.3%; SU
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                               0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19
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                                                                                                                                                                                                                                           0;
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Pred. No.
                                                                                                                                                                                                                                                                       Score 15.8;
Pred. No. 7
                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 using specified oligo-nucleotide
                                                                                                                                                                                                                                                                  .7e+02;
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                                                                                                                                                                                                                                                                                           Length
                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                                                                                                                                           <u>.</u>
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                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                           0
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AAT30413 standard; DNA; 19 BP

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RESULT 535
ADF31861
ID ADF318
XX ADF318
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect pd. simple sequence repeat (SSR), and an adaptor directed primer, comprising a sequence complementary to an adaptor segment. The present sequence is an example of a compound SSR primer. The method represents a modified amplified fragment length polymorphism assay, which is partic. useful for genome fingerprinting, i.e. for genetic trait marking and germplasm comparisons
             RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; proliferative disease; restenosis; polycystic kidney disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellice region; genetic trait marking; germplasm comparisons; compound; ss.
                                                                                                                                                     Human IGF-1R siNA lower strand,
                                                                                                                                                                                        12-FEB-2004
                                                                                                                                                                                                                             ADF31861
                                                                                                                                                                                                                                                                 ADF31861 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 9 A; 0 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 2; Page 84; 173pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modified amplified fragment length polymorphism assay - for detection polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Morgante M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            28-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Compound simple sequence repeat primer (GA) 7.5 (TA) 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-JAN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT30413;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9617082-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (DUPO ) DU PONT DE NEMOURS & CO
 Inflammatory disease; allergic disease;
                                                                                                                                                                                                                                                                                                                                                                                               1183 GAAAGAGAGAGAGAAAT 1201
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1996-277795/28.
                                                                                                                                                                                                                                                                                                                                                             GAGAGAGAGAGAGATAT 19
                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94US-00346456
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
                                                                                                                                                                                                                                                                 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.8; DB 1; Pred. No. 7.7e+02;
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                                                                                                                                                     SEQ ID NO:526
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 19;
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RESULT 536
ADF31584/c
ID ADF315
XX
AC ADF315
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                                                                                                                                                                                                                                                                                   CC further comprise sense and antisense regions, or alternatively are consembled from a sense oligonucleotide and an antisense oligonucleotide. CC Specifically, the siNAs include short interfering RNA (siRNA), double-cC stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs CC can be unmodified or chemically modified, can contain CC deoxyribonucleotides, and can be chemically synthesised, expressed from a CC vector or enzymatically synthesised. The invention also relates to kits CC for the in vitro or in vivo delivery of siNA; conjugates and/or complexes CC expression of the IGF-IR gene in cells, tissue explants or organisms CC (e.g., by ex vivo gene therapy), or in grafts and transplants for the CC treatment of a variety of conditions. They may be used for treating CC cancer and other proliferative diseases (e.g., restenosis and polycystic CC kidney disease), inflammatory and/or allergic diseases, autoimmune CC diseases and transplant rejection. The siNAs are also useful for drug creening, diagnosis, therapeutic target identification and validation, CC genetic engineering, pharmacogenomics, studying gene function, and gene calping (e.g., of single nucleotide polymorphisms). The present sequence CC represents the lower strand of a human IGF-IR-targeted double-stranded
                                                                                                                                                                                                         Query Match
Best Local S
Matches 17
                                      ADF31584 standard; RNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-MAR-2002;
06-JUN-2002;
     ADF31584
                                                                                                                                                                                                                                                                                 Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to short interfering nucleic acids (siNA) which downregulate expression of the human insulin-like growth factor 1 receptor (IGF-IR) gene by RNA interference. The siNAs may or may not comprise ribonucleotides and may be double or single stranded. They
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the insulin-like growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-721691/68.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-SEP-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2003; 2003WO-US005044.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transplant rejection; cytostatic; vasotropic; nephrotropic; antiinflammatory; antiallergic; immunosuppressive; human; insulin-like growth factor 1 receptor; IGF-1R; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JAN-2003;
                                                                                                                                                                    2568 GGAGAGAGAGATGGAGAAC 2586
                                                                                                                                   _
                                                                                                                                                                                                           17;
                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               receptor gene.
                                                                                                                                   GGAGAGAGAAAGGGAGAAC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 526; 147pp; English.
                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                 BP; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2002US-0406784P.
2002US-0408378P.
2002US-0409293P.
2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2002US-0363124P.
2002US-0386782P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002US-0358580P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Beigelman L,
                                                                                                                                                                                                                                                                                 A; 1 C; 9
                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                         <u>,</u>
                                                                                                                                                                                                                                                                                 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chowrira
                                                                                                                                                                                                                         Score 15.8; DB 1
Pred. No. 7.7e+02
                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            В
                                                                                                                                                                                                                                            DB 1; Length 19;
                                                                                                                                                                                                           Indels
                                                                                                                                                                                                           0
                                                                                                                                                                                                         Gaps
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comprise ribonucleotides and may be double or single stranded. They comprise ribonucleotides and may be double or single stranded. They comprise ribonucleotides and may be double or single stranded. They comprise ribonucleotides and antisense regions, or alternatively are further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the siNAs include short interfering RNA (siRNA), doublestranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be unmodified or chemically modified, can contain deoxyribonucleotides, and can be chemically synthesised, can contain deoxyribonucleotides, and can be chemically synthesised from a cyector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA, and vectors that express siNA. The siNAs are used to modulate expression of the IGP-IR gene in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for treating therapy in the sinas are used to modulate the cancer and other proliferative diseases (e.g., restenosis and polveystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409293P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; mi short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; cancer; proliferative disease; restenosis; polycystic kidney disease; inflammatory disease; restenosis; polycystic kidney disease; transplant rejection; cytostatio; vasotropic; nephrotropic; antiinflammatory; antiallergic; immunosuppressive; human; insulin-like growth factor 1 receptor; IGF-1R; target sequence;
                                      cancer and other proliferative diseases (e.g., restenosis and polycystic kidney disease), inflammatory and/or allergic diseases, autoimmune diseases and transplant rejection. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the upper strand of a human IGF-IR-targeted double-stranded siNA, which is identical to the IGF-IR transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates expression of the insulin-like growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mcswiggen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human IGF-1R transcript target sequence/siNA upper strand, SEQ ID NO:249.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to short interfering nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 3; SEQ ID NO 249; 147pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-FEB-2003; 2003WO-US005044.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ų,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Beigelman
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chowrira B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              micro-RNA; miRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (siNA) which
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Query Match Best Local Similarity

0.3%;

Score 15.8; DB 1; Pred. No. 7.7e+02;

Length 19;

Sequence 19

BP; 0 A; 9 C; 1 G; 0 T; 9 U; 0 Other;

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RESULT 537
AAQ75581/c
ID AAQ755
XX JP063C
XX AAALY
PT AAALY
CCC COUDL
CCC COUDL
CCC COUDL
CCC COUDL
CCC COUL
AAT41429
ID AAT4144
XX AAT414
XX AAT414
XX O2-FEB
DT 02-FEB
DX Rat ob
XX Obese
XX Synthe
XX Synthe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type (abbelled reverse transcription primers (CENESEO files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                           Synthetic
                                                                              Obese gene;
                                                                                                                                                                                     02-FEB-1997
                                                                                                                                                                                                                                         AAT41429
                                                                                                                                                                                                                                                                                          AAT41429 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 0 C; 0 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         method can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 5; 11pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analysis of cDNA and gene expression - by by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          aggregate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Analysis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ75581 standard; DNA;
                                                                                                                               Rat obese gene antisense primer No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NITS ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19
                                                                                                                                                                                                                                                                                                                                                                                                                            20
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                                                                                                                                                                                                                                                                                                                                                                                                                            ATTAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTAAAAAAATACAAAAAA 5408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGAGAGAGAGATGGAGAAC 2586
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                restriction
                                                                              မှ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                   (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93JP-00112515
                                                                              gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93JP-00112515
                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
                                                                              obesity; polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             enzyme;
                                                                                                                                                                                                                                                                                             20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.8; DB 1; Length Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                  7.
                                                                              chain reaction; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                              primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 539
AAA55551/c
ID AAA555
XX AAA555
XX JO-AUG
DT JO-AUG
DT TRAF2
XX TUMOUZ
KW TUMOUZ
KW ANTi-1
XX Synthe
XX WO2000
PD 13-APF
XX 06-OCT
XX 06-OCT
XX USIS-
XX ANTISE
PT NECTOS
PT NEC
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primers 5-8 (AAT41427-30) are based on a 184 bp rat obese gene cDNA clone isolated from epididymis fat tissue (see also AAT41426). Primers 5 and 7, and primers 6 and 8, were used in 2 PCR reactions to amplify cDNA derived from epididymis fat tissues of Sprague- Dawley rats. Amplified products were cloned into pBluescript, introduced into E. coli JM109 and sequenced. A complete rat obese gene cDNA (AAT41421) was obtd
                                                                        Antisense oligonucleotides targeted to nucleic acids encoding human tumor necrosis factor receptor-associated factor (TRAF), useful for treating diseases associated with TRAF expression such as inflammatory diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-APR-1996;
                                                                                                                                                                                     WPI; 2000-303732/26
                                                                                                                                                                                                                                     Baker BF,
                                                                                                                                                                                                                                                                                                                                         06-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                          05-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                             13-APR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200020435-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    anti-inflammatory; B-selectin; jun kinase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tumour necrosis factor receptor-associated factor; TRAP; human; antisense oligonucleotide; phosphorothioate; antiproliferative;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TRAF2 antisense oligonucleotide ISIS# 16842.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-AUG-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA55551;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA55551 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 6 A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 7; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-444886/45
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-APR-1995;
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                                                                                                                                                                                                                                                                                      (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             recombinant rat obese gene - used to develop prods. to study obesity to diagnose obesity and obesity factors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ζ,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TGAGGATACCTGGGAGCCA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ogawa Y,
                                                                                                                                                                                                                                     Cowsert LM,
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Pred. No. 7.7e+02
0; Mismatches
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Example 16; Page 52; 170pp; English

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RESULT 540
AAA558
ID AAA558
XX AAA558
XX AAA558
XX Human;
XW Human;
XW Human;
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XY HOMO 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; DNA methyltransferase; DNA MeTase; antisense oligonucleotide; modulation; inhibition; gene expression; combination therapy; p16; histone deacetylase; HDAC; thymidylate synthase; tumour suppressor; methylation; gene therapy; tumour; cytostatic; antiasthmatic; antiinflammatory; inflammation; asthma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            selectin expression in cells or tissues comprises contacting the cells of tissues with an antisense oligonucleotide targeted to TRAF-6. The antisense oligonucleotides have antiproliferative and anti-inflammatory activity and are useful for treating disorders associated with cell proliferation and inflammation. The antisense oligonucleotides
                                                                                                                                                                                                                                                                               Inhibiting gene expression e.g. DNA methyltransferase, with a synergistic amount of antisense oligonucleotide effectors e.g. 5-aza-cytidine of gene products, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-339532/29.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human histone deacetylase HD2 antisense oligonucleotide SEQ ID NO:51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA55806
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA55806 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (METH-) METHYLGENE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to antisense oligonucleotides (see AAA55496-
A55757) which are targeted to nucleic acids encoding a human tumour
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
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No. 7
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                                                                                                                                                                                                                                                                                   by treating cells and protein for gene therapy
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The present invention describes a method for inhibiting the expression a gene in a cell comprising contacting the cell with an effective synergistic amount of an antisense oligonucleotide which inhibits expression of the gene, and an effective synergistic amount of a protein

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Primers AAZ32974-Z32977

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encoding

Disclosure; Page 29; 99pp; English.

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RESULT 541
AAZ32975
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Best Local Similarity
Thes 17; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC effector of a product of the gene. Also described are: (1) a method for CC treating a disease responsive to inhibition of a gene in a mammal; (2) a CC method for inhibiting tumour growth in mammal; (3) an inhibitor of a gene CC comprising an antisense oligonucleotide which inhibits expression of the CC gene in operable association with a protein effector of a gene product; CC and (4) a pharmaceutical composition comprising the inhibitor of (3). The CC methods and compositions are useful as analytical tools for transgenic CC diseases including benign and malignant tumours, inflammation or asthma. CC expression or activity of a gene or gene product may be used to treat CC expression or activity of a gene or gene product may be used to treat CC patients having, or predisposed to developing, a disease responsive to CC inhibition of the gene. These may also be used to activate silenced genes to provide missing gene functions and improve a given condition. CC Physiological function of a gene product in an experimental cell culture or appression. ARAS5758 to ARAS5842 represent of inhibiting gene activity invertion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
                             Claim 8; Col 33-34; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human MKK4 exon A PCR primer CG2exA.RP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAZ32975 standard; DNA;
                                                                         Polynucleotides comprising all or a MKK4 locus are useful for diagnosis,
                                                                                                                                                       Skolnick MH,
                                                                                                                                                                                                                     10-JAN-1997;
                                                                                                                                                                                                                                                  13-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                        somatic; signal transduction; apoptosis; stress; cytokine; induction; phosphorylation; Jun kinase; JNK; p38; tumour; suppressor; loss of heterozygosity; LOH; cancer; detection; diagnosis; prognosis; breast cancer; pancreatic cancer; coloer; testicular cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MKK4; mitogen activated protein kinase; MAPK; MAPK pathway;
                                                                                                                                                                                      (MYRI-) MYRIAD GENETICS INC.
                                                                                                                                                                                                                                                                                 23-NOV-1999
                                                                                                                                                                                                                                                                                                              US5989885-A
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                                                                                                                                                                                                                                                                                                                                                                                                           grug
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                                                                                                                                                                                                                                                                                                                                                                                                          screening;
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                                                                                                                                                     Perry WL,
                                                                                                                                                                                                                     97US-00782482.
                                                                                                                                                                                                                                                  97US-00874186.
                                                                                                                                                                                                                                                                                                                                                                                                          gene therapy; protein replacement therapy; mimetic; PCR
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Pred. No. 7.
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                                                                          portion of the , prognosis and
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                                                                          therapy of human
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WPI; 2000-664982/64.

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ARC79540
ID AAC7957
XX AAC795
XX AAC795
XX AAC795
XX AAC10
DE Murine
XX Antirh
KW Antirh
KW Antirh
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XX AAC10
DE MUS 8P
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antisense oligonucleotide; p38 mitogen activated protein kinase; MAPK; antirheumatic; antiarthritic; immunosuppressive; cardiant; heart disease; antiinflammatory; autoimmune disease; rheumatoid arthritis; apoptosis; phosphorothioate; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                       WO200059919-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murine p38beta antisense oligonucleotide SEQ ID 65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-FEB-2001 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAC79540;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAC79540 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 6 A; 1 C; 13 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           function for cancer therapy. MKK4 gene therapy, protein replacement therapy and protein mimetics that reconstitute the function of the Mi protein may be used for therapy of human cancers which result from a
                                       Monia BP,
                                                                                                                                                                                                                                                                     04-APR-2000, 2000WO-US008794
                                                                                                                                                                                                                                                                                                                                                12-OCT-2000
                                                                                                                  (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2566 GGGGAGAGAGAGATGGAGA 2584
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGGGAGGAGAGAGGGAGA 20
                                       Gaarde WA, Nero
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                         9908-00286904
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                                       PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.8; DB 1; Length 20; Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                       Mckay R,
                                   Popoff I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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This invention relates to antisense compounds 8-30 nucleobases in length CC targeted to the 5'-untranslated region, translational start site, CC translational termination region of 3'-untranslated region of a nucleic acid encoding a p38 mitogen activated protein kinase (MAPK), where the antisense oligonucleotides inhibit the expression of MAPK. Sequences AAC79480 and AAC79501 represent human p38alpha MAPK and p38beta MAPK cDNA cequences. AAC79571 - AAC79500 and AAC79553 - AAC79570 represent human p38beta antisense oligonucleotides, while AAC79502 - AAC79571 and CC AAC79571 - AAC79580 represent human p38beta antisense oligonucleotides. Also included in the invention are a p38alpha cDNA sequence AAC79523 and antisense oligonucleotides AAC79523 - AAC79536 isolated from rat tissue. The antisense oligonucleotides have antirheumatic; antiarthritic; The antisense oligonucleotides have antirheumatic; antiarthritic; CC oligonucleotides are useful for inhibiting the expression of p38 MAPK in CC oligonucleotides are useful for inhibiting the expression of p38 MAPK in CC oligonucleotides are useful for inhibiting the expression of p38 MAPK in CC oligonucleotides are useful for inhibiting the expression of p38 MAPK in C oligonucleotides are useful for inhibiting the expression of p38 MAPK in C oligonucleotides are used for tracting an animal CC with disease such as inflammatory or autoimmune diseases e.g. rheumatoid arthriting of heart disease. The oligonucleotides are also useful for inhibiting the expression of p38 MAPK in C inhibiting the expression of p38 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense compound targeted to p38 mitogen activated protein kinase inhibits protein kinase and is useful for diagnosing and treating inflammatory, autoimmune and heart disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 5; Page 53; 90pp; English.
                                                                               inhibiting inflammation or apoptosis
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밁 Ś Query Match
Best Local Similarity
Matches 17; Conserv 2639 CCCTGCAGCTGCTGCTGCA 2657 1 CCCTGCAGCCGCTGCGGCA 19 Conservative 0.3%; 0; Mismatches Score 15.8; DB 1 Pred. No. 7.7e+02 DB 1; Length 20; Indels 0 Gaps 0

Sequence 20 BP; 2 A; 10 C; 6 G; 2 T; 0 U; 0 Other;

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RESULT 543
AAH43116
ID AAH431
XX AAH431
XX AAH431
XX AAH431
XX AAH18
AX AAT18

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Antisense oligo, target HDAC-2 121-141.
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WO200138322-A1. Synthetic.

31-MAY-2001

22-NOV-2000; 2000WO-IB001881.

23-NOV-1999; 99US-0167035P

(METH-) METHYLGENE INC.

Delorme D, Ruel R, Lavoie R, Thibault C, Abou-Khalil

2001-432601/46.

New inhibitors of histone deacetylase e.g. N-hydroxy-5-(4-(benzenesulfonylamino)-phenyl)-4-yn-2-pentanamide for trearestenosis or fungal infections. treating cancer,

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RESULT 544
AAC81378
ID AAC813
XX AAC813
XX AAC813
XX AAC813
XX AAC813
XX AAC813
XX VIVANA
Cell B
XX VIVANA
XX CANCEY
XX CANCEY
XX Phomo B
XX VIVANA
XX AAC8140
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Sequences AAC81326-C81405 represent antisense oligonucleotides targetted to the human Y-box binding protein 1 gene, which inhibit its expression. The antisense oligonucleotides were designed to target different regions of the human Y-box binding protein 1 mRNA, and were analysed for their effect on Y-box binding protein 1 mRNA levels by quantitative real-time PCR. Human Y-box binding protein 1 (also known as YB-1, DNA binding protein B and dbpB) is a member of the Y-box binding protein factors, a highly conserved family of nucleic acid binding proteins which bind to the Y-box, an inverted CCAAT sequence found in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The sequences given in AAH43115-21 are oligonucleotides which are antisense to the histone deacetylase gene, HDAC-2. These oligonucleotides may be used in combination with an inhibitor of histone deacetylase enzyme function, to given an improved inhibitory effect, thereby reducing the amount of inhibitor required to obtain a given inhibitory effect. Compounds containing these oligonucleotides may be used to treat cell proliferation conditions such as cancer, restenosis or psoriasis. They can also be used to treat protozoal and fungal infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human Y-box binding protein 1; YB-1; DNA binding protein B; dbpB; transcription factor; nucleic acid binding; DNA repair; cell sensitiaation; genotoxic stress; immune regulation; MHC expression; viral gene expression; extracellular matrix degradation regulator; redox signalling; expression inhibition; tumour formation; cancer multidrug resistance; inflammation; immune disorder; infection;
                                                                                                                                                                                                                                                                                                                                                                                                              Antisense oligonucleotides, useful for modulating the expression of Y-box binding protein 1, as well as for treating or preventing diseases associated with Y-box binding protein 1 expression, e.g. inflammation or
                                                                                                                                                                                                                                                                                                                         Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human Y-box binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-023284/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 40; 147pp; English
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                                                                                                                                                                                                                                                                                                                                                                                          formation.
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                                                                                                                                                                                                                                                                                                                         Col 43-44; 40pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          99US-00429323.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein 1 antisense oligonucleotide, SEQ ID NO:62.
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No. 7
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RASSULT 545
AAC89545
ID AAC895
XX AAC895
AC AAC895
XX AAC895
XX AAC895
XX OB-MAR
XX OB-MAR
XX Histon
KW HDAC-I
XX HOMO 6
XX HO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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Best Local (
    Sequence 20
                                                                                                                           The present invention provides inhibitors of histone deacetylase enzymenesuch as HDAC-1, HDAC-2, HDAC-3, HDAC-4, HDAC-5, HDAC-C and HDAC-D. These inhibitors may be antisense strands or they may be compounds identified by contacting the enzyme with the compound and measuring the resulting enzyme activity. These inhibitors are useful for treating cancers and for example of the contacting the enzyme with the compound and measuring the resulting enzyme activity. These inhibitors are useful for treating cancers and for the contact in the contact
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense oligonucleotide that inhibits expression of a histone deacetylase, useful for treating and/or alleviating the symptom neoplasia, or for inhibiting neoplastic cell growth in an anima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  gene therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Macleod AR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-MAY-2000; 2000WO-IB001252
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             08-MAR-2001
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                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 24; 125pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99US-0132287P
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             G; 4 T; 2 U; 0 Other;
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Query Match Best Local Similarity

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Score Pred.

15.8; DB 1; No. 7.7e+02;

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RESULT 546
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Matches 17; Conserv
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deacetylase, us
                                                                                                                                                                                                                                                                                                                                                                         The present invention provides inhibitors of histone deacetylase enzymes such as HDAC-1, HDAC-2, HDAC-3, HDAC-4, HDAC-5, HDAC-C and HDAC-D. These inhibitors may be antisense strandacrotthey may be compound identified by contacting the enzyme with the compound and measuring the resulting enzyme activity. These inhibitors are useful for treating cancers and for identifying which histone deacetylase is involved in a neoplasia
                                                            09-SEP-2004
07-SEP-2001
                                                                                                           AAS05714;
                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure, Page 12, 125pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Antisense oligonucleotide that inhibits expression of a histone deacetylase, useful for treating and/or alleviating the symptoms neoplasia, or for inhibiting neoplastic cell growth in an animal.
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reverse phase triplex forming oligonucleotide; RP-TFO;
                                                                                                                                        AAS05714 standard;
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                             Aminopurine substituted region of an RP-TFO.
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                                                            (revised)
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Pred. No. 7.7e+02;
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                                                                                                                                                                             associated oncogene-1, for diseases associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; glioma-associated oncogene-1 associated disease; infection, inflammation; tumour formation; cytostatic; antiinflammatory; anti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK30536 standard;
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                                                                                                                        Claim 1;
                                                                                                                                                                                                                                                                                                         WPI; 2002-138363/18.
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                                                                                                                                                                             antisense compounds targeted to nucleic acids encoding glioma-
liated oncogene-1, for modulating the gene expression and treati-
ses associated with expression of the oncogene in humans.
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                                                                                                                                                                                                                                                                                                                                                                 Wyatt J;
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The present invention relates to antisense compounds and methods modulating the expression of human glioma-associated oncogene-1.

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RESULT 549
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The present invention relates to novel human nucleic acid sequences (I). The present sequence is one such nucleic acid sequence. Expression of (I) are implicated in tumour suppression or reversion and apoptosis and viral resistance. (I) are useful as probes or primers for detecting, identifying, measuring and/or amplifying nucleic acid sequences, as antisense reagents and for recombinant production of polypeptides. (I), polypeptides (II) encoded by (I), vector containing (I), cells containing these vectors and antibodies (Ab) against (II) are all useful for treatment/prevention of viral, tumour and cell degeneration diseases (especially neurodegeneration, such as Alzheimer's disease and schizophrenia). Analysing the expression of (I) is also useful for diagnosis and/or prognosis of such diseases. Transgenic animals carrying (I) are used for studying the actiology of these diseases (also immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New nucleic acid implicated e.g. in diagnosis of tumors, viral infection
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tion and cellular degeneration and
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RESULT 550
ABL4554
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DT 11-APF
DE Human
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                                                                                                                                                  (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell plates containing the clones having said marker sequence; (d) the order of the markers is changed so that the same discrimination Nos. succeed to the maximum in the specified discrimination Nos. to array the multiwell plates (e) the clones in the multiwell plates of the specified discrimination Nos. are mixed respectively in each wells of longitudinal and lateral directions; (f) the mixed clones are cultured and the resultant cultures are amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (1) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL42572 to ABL45322 represent PCR primers for human chromosome 21q22.1, which are specifically claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 17
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention describes a method of arraying genome clones. The method comprises: (a) clones of the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human chromosome 21q22.1 PCR primer SEQ ID NO:2590.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and inflammatory diseases). Note: In the present specification, SEQ ID 1 to 2280 are claimed in Claim 1, however only SEQ ID 1 to 2270 are shown in the specification
                                                                                             Sequence 20 BP; 4 A; 6 C; 4 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 6; Page 56; 528pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Arraying genome clones
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (RIKA ) RIKAGAKU KENKYUSHO. (GENO-) GENOTEX YG.
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Score 15.8; DB 1; Pred. No. 7.7e+02;
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Pred. No. 7.
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7.7e+02;
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RESULT 551
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WPI; 2002-416486/44
                                          Ward DT,
                                                                                   (ISIS-) ISIS PHARM INC
                                                                                                                              29-SEP-2000; 2000US-00676436.
                                                                                                                                                                       28-SEP-2001; 2001WO-US030549.
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                                          Monia BP,
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                                          Wyatt JR;
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RESULT 552
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XX ABX046
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New antisense compound targeted to activated protein kinase 4, useful inflammatory disorder or cancer.
  The invention describes novel isolated polypeptides (I, Ib) useful for diagnosing prostate cancer comprising obtaining a patient sample containing prostate cells and detecting the presence or absence of an expression product of a HML-2 endogenous retrovirus in a patient sample. Polynucleotides associated with (I) are useful for diagnosis or treatment of testicular cancer, multiple sclerosis or insulin-dependent diabetes mellitus. An inhibitor of a HML-2 protease and a transdominant negative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              diagnostics,
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07-DEC-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          cancer; transgenic animal; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; endogenous retrovirus; herv; prostate cancer; testicular cancer; multiple sclerosis; insulin-dependent diabetes mellitus; HML-2 protease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human endogenous retrovirus k (herv-k) associated probe #27.
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The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary coligonucleotide probes (II) will hybridise with little mismatch, where coligonucleotide probes (II) will hybridise with little mismatch, where coligonucleotide probes (II) will hybridise with little mismatch, where coligonucleotide probes (II) which is apported to the method is useful for detecting infectious diseases caused by bacterial infectious agents colifications and haemophilus influenza, fungal confectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus funigautus, viruses e.g. T-cell lymphocytotrophis cirus, compared to the colification and polio virus, and parasitic infectious agents selected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, psi gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food
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                       The present invention describes a method (M1) for designing capture oligonucleotide probes (II) wall hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents e.g. Cryptococcus neoformans, Candida albicans and infectious agents e.g. Cryptococcus neoformans, Candida albicans and Aspergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and polio virus, and parasitic infectious agents escapetions. Epstein-Barr virus and polio virus, and parasitic infectious agents medinesis. The method is also useful for detecting genetic diseases such as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. Detecting cancer involving oncogenes, tumour suppressor genes, or genes cancer is specifically associated with a gene selected from BRCA1 gene, parasitic and the food action of the food and it also useful for for detecting cancer. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and feed industry, detecting comprises scanning (using e.g. a scanning electron microscope and infrared microscope) the support at the particular sites and identifying if ligation of the oligonucleotide probe sets occurred and correlating (using a computer) identified ligation to a presence or absence of the target nucleotide sequences. ABI82074 to ABI97546 represent oligonucleotide sequences used in the exemplification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Designing capture oligonucleotide probes complementary oligonucleotides hybridize
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-APR-2001; 2001WO-US010958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Capture oligonucleptide Zip ID#2084 oligo
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fig 29; 300pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP;
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Pred. No. 7.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                              Monia BP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   06-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-SEP-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense; antiarthritic; antiinflammatory; kinase inhibitor; inflammatory disease; rheumatoid arthritis; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             p38 mitogen-activated protein kinase; p38 MAPK; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murine p38-alpha MAPK antisense oligonucleotide ISIS NO 100802
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                                                                                                                                                                                                                                                                                                                      New antisense compound,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note- "nucleotides 1-5 & 16-20 are 2'-methoxyethoxy (MOB) nucleotides, nucleotides 1-4 & 16-19 are 11x via phosphodiester linkages, nucleotides 6-15 are 2'deoxy- nucleotides, nucleotides 5-16 are linked via phosphorothioate linkages, all C nucleotides are 5-
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Pred.
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                                                                                                                                                                                                                                                                                           preparing a composition for inflammatory diseases, e.g.
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No. 7.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     kinase inhibitor; mouse;
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This invention describes a novel antisense compound, which is 8-30 nucleobases in length targeted to a nucleic acid molecule encoding p38 mitogen-activated protein kinase (MAPK). The products of the invention have antiarthritic and antiinflammatory activity, can act as act as kinase inhibitors. The antisense compound is useful for preparing a composition for diagnosing, treating or preventing inflammatory diseases, e.g. rheumatoid arthritis or for use in antisense gene therapy. This

Example 5;

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27-28; 44pp;

English.

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Best Local
                                                                                                                                                                                                                                                                The invention relates to a novel support for ligand immobilization, which is coated with a polyanion which has previously been activated. The support is useful for the production of DNA chips, protein chips, sugar chips and biosensors for investigative and diagnostic uses. Ligands which can be immobilized to the support include agonists, antagonists, toxins, venoms, virus epitopes, hormones, lectins, hormone receptors, peptides, nucleic acids, drugs, sugars, oligonucleotides, proteins, antigens, monoclonal antibodies, cells, viruses, and avidins. In an example of the invention, the ligand bound to the support is a PCR primer targeted to a number of genes and used to diagnose the presence and potentially the transcription of the genes. This sequence represents a 5' primer targeted to the v-akt murine thymoma viral oncogene homologue 1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ACC44257 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 33; 51pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polyanion-coated ligand immobilization support for production of DNA chips, protein chips and biosensors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-342750/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Asada K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA chip;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5' primer to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-JUL-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACC44257;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 2 A; 10 C; 6 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence represents an inhibit p38 MAPK
                                                                                                                                                                                                                     Sequence 20 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-SEP-2001; 2001JP-00288149
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20-SEP-2002; 2002WO-JP009661.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer; ss; support; ligand immobilization; activated polyanion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (TAKA-)
                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2639 CCCTGCAGCTGCTGCA 2657
                                                   5167 TGTCACAGGGTGAGCCCCA 5185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17;
19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAKARA BIO INC.
                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CCCTGCAGCCGCTGCGGCA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Imose N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein chip;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amplify v-akt homologue gene for ligand support method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                               A; 7 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Takeda 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                 0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense oligonucleotide used in a method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sugar chip; biosensor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
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                                                                                                       0;
                                                                                                                              Pred.
                                                                                                                                          Score 15.8;
Pred. No. 7.
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Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rokushima
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                                                                                                                                    7e+02
                                                                                                                                                              DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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RESULT 557

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ACC70061/c
ID ACC700
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ACC ACC700
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PG PG PT
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PG PG PT
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PG PG PT
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NO2003
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ON-MAY
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INFLAM
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IN
   RESULT 558
ACC85486/c
ID ACC854
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AC ACC854
XY
DT 29-SEP
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DE Human
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KW Late P
KW exces8
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primers ACC70061-62 were used to amplify cDNA encoding a pig corticosteroid-binding globulin (Cbg). The specification describes a method of identifying polymorphic markers associated with the hypercortisolemia phenotype. The method comprises comparing nucleic acid sequences, from many individuals, that contain at least part of the Cbg gene and identifying mutations in this gene or adjacent sequences. The method is used for diagnosis of hypercortisolemia, or predisposition to it, for identifying dysfunction of the corticotropic acid and associated diseases, e.g. obesity; constitutive sensitivity to inflammatory or autoimmune reactions; aging (especially cognitive) or sensitivity to illicit drugs, It is also useful for (counter-) selection of breeding animals, especially pigs, with a high probability of developing hypercortisolemia and obesity
                                                                                           Human HEX-A gene PCR primer SEQ ID NO: 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACC70061;
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                                                                                                                                                                                                                                                                               ACC85486 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying polymorphic markers associated with hypercortisolemia, useful for diagnosis of dysfunction of the corticotropic axis and for selection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-430536/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-OCT-2001; 2001FR-00014156.
26-JUL-2002; 2002FR-00009551.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-OCT-2002; 2002WO-FR003762.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        08-MAY-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JUL-2003
Late PCR; polymerase chain reaction; human; CFTR; HEX-A; limiting primer; excess primer; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 16; 50pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of breeding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Moisan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pig; corticosteroid-binding globulin; Cbg gene; polymorphic marker;
hypercortisolemia; corticotropic acid; obesity; constitutive sensitivity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR primer for
                                                                                                                                                       29-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            652 CAGCCAGAGAAGACGAGTG 670
                                                                                                                                                                                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                   CAGCCAGAGAGACAGGTG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mormede
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 2 A; 7 C; 4 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Milan D,
                                                                                                                                                                                                                                                                                  ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 7.76
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.8; DB 1;
Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bidanel J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ousova
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
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RESULT 559
AL61584/c
ID AAL615
XX AAL615
XX AAL615
XX Human
XX Human;
KW Human;
KW Ikappa
KW Ikappa
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XX Homo s
OS Synthe
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a method of non-symmetric polymerase chain reaction (PCR) amplification, comprising thermally cycling a PCR reaction mixture containing a DNA amplification target sequence, a pair of PCR primers, dNTP's and thermostable polymerase repeatedly through PCR steps of strand melting, primer annealing, and primer extension. The method is useful for amplifying stretches of DNA, including cDNA reverse transcribed from RNA, for assays, for diagnostics and other purposes. The present sequence is a primer used to demonstrate the method of the
                                                                                                                                                                                                                Human; inhibitor-kappa B-R; I-kappaBR; IKBR; I-kappa-B-related; NFKBIL2; ikappab r; antisense; immune response; infection; inflammation; therapy; tumour; prophylaxis; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Non-symmetric polymerase chain reaction amplification, useful for amplifying DNA by thermally cycling a PCR reaction mixture containing DNA amplification target sequence, a pair of PCR primers, dNTP's and
                                                                                                                                                                                                                                                                      Human inhibitor-kappa B-R antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 66; 125pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-DEC-2001; 2001US-0341886P
17-DEC-2002; 2002US-00320893
                                                              modified_base
                                                                                                                               modified_base
                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                   22-SEP-2003
                                                                                                                                                                                                                                                                                                                              AAL61584
                                                                                                                                                                                                                                                                                                                                                        AAL61584 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               thermostable polymerase.
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                                                                                                                                                                                        Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 A;
/*tag= b
/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                            /mod_base= OTHER
/note= "Phosphorothioate
are 5-methylcytidines"
                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 C; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.8;
Pred. No. 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .7e+02;
                                                                                          backbone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            J, Sanchez
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                          All cytidine residues
                                                                                                                                                                                                                                                                        ISIS #130509.
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RESULT 560
ADG25637
ID ADG256
XX ADG256
XX ADG256
XX TObacc
XX Plant;
KW plant;
KW plant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to antisense compounds targetted to a nucleic acid molecule encoding human inhibitor-kappa B-R (also known as I-kappaBR, I-Kappa-B-related, ikappab r, nuclear factor of kappa light polypeptides gene enhancer in B-cells inhibitor-like 2 and NFKBILD) to inhibit its expression. Antisense compounds of the invention are useful for treating diseases or conditions associated with the expression of inhibitor-kappa B-R such as a heightened immune response involving increased cytokine expression, or a result of infection (e.g. bacterial, viral or parasitic). They are useful for diagnostics, therapeutics, prophylaxis e.g. to prevent or delay infection, inflammation or tumour formation, as research reagents and kits and in distinguishing between functions of various members of a biological pathway. They are also useful in antisense therapy. The present sequence is an oligonucleotide targetted to human inhibitor-kappa B-R DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                         plant; herbicide; serine hydroxymethyltransferase; SHMT: plant growth regulator; desiccant; defoliant; herbicide transgenic plant; mitochondrial; ss; primer; PCR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides targeted to nucleic acids encoding inhibitor-kappa B-R, useful for diagnosing or treating diseases associated with expression of inhibitor-kappa B-R, e.g., a heighte immune response or infection.
   13-MAR-2003; 2003WO-EP002574.
                                                                          WO2003078613-A2
                                                                                                            Nicotiana tabacum
                                                                                                                                                                                                                                                     26-FEB-2004
                                                                                                                                                                                                                                                                                         ADG25637;
                                                                                                                                                                                                                                                                                                                            ADG25637 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 3 A; 7 C; 7 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3; Page 75; 108pp; English.
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                                                                                                                                                                                                                SHMT PCR primer SHMT1*2
                                                                                                                                                                                                                                                                                                                                                                                                                    TGCAGCTGAGGCTGCAGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                     (first entry)
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/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
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                                                                                                                                                                  resistance;
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RESULT 561
ABZ88038/c
ID ABZ880
XX ABZ880
XX ABZ880
XX Human
XX Human;
KW Human;
KW antias
KW antias
KW lung i
XX Homo E
XX Ho
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Best Local S
Matches 17
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Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
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                                                                                       WPI; 2003-229219/22.
                                                                                                                                         Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200285308-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammation;
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                                                                                                                                   Li Y,
Tang
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Reindl A, Schmidt
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                                                                                                                                         Sandrasagra A,
L, Shahabuddin
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0; Mismatches
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S;
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corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                        ubiquinone
                                                                                  SEQ ID
                                                                                 NO 3280; 872pp; English
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initiation codon, coding region, 5' or 3' end genomic flanking regions, c5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or masal airway dysfunction and a second active agent comprising an antiinflammatory, antiallergic, antiasthmatic, hypotensive, has antiinflammatory, antiallergic, antiasthmatic, hypotensive, and cytostatic activity. The composition may have a composition is useful for treating or greventing a respiratory, lung or malignant disease or condition, also cfor enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing depleting levels of, or reducing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung surfactant in a subject's tissue, or treating bronchoconstriction, condition, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO specification, but was obtained in electron:
at ftp.wipo.int/pub/published_pct_sequences The invention relates to active agent comprising an a novel pharmaceutical composition, which has ising an oligonucleotide antisense to the region, 5' or 3' end genomic flanking regions,

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                                   Query Match
Best Local S
Matches 17
                                                                      Sequence 20
                  2641 CTGCAGCTGCTGCAGC 2659
                                   l Similarity
17; Conserv
                                                                       BP; 3 A; 7 C; 9 G; 1 T; 0 U; 0 Other;
                                    Conservative
                                           0.3%;
                                    0;
                                             Score 15.8;
Pred. No. 7.
                                    Mismatches
                                  7.7e+02;
2;
                                                      DB 1; Length 20;
                                     Indels
                                    0
                                    Gaps
                                     0
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RESULT 562
                                                              Human oligonucleotide sequence.
                                                                  17-OCT-2003
                                                                       ABZ85534
                                                                            ABZ85534 standard;
                                                                  (first
                                                                            DNA;
                                                                            20
                                                                            BP.
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Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antianflammatory steroid; ubiquinone; antianflammatory; can therapy; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; respiratory; lung; adenosine sensitivity; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; Homo sapiens. inflammation; respiratory disease; ds.

WO200285308-A2

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC.

Li Y, Tang Ļ, Sandrasagra Shahabuddin Þ ŝ Katz (F) Pabalan ŗ,

2003-229219/22.

Pharmaceutical composition respiration, has oligo(s) a n for treating ailments associated with impaired antisense to specific gene(s) or its

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RESULT 563
ABZ90836
ID ABZ908
XX ABZ908
XX ABZ908
XX Human
DE Human
XX Homo (
XX ADZ002
XX Homo (
XX Hom
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Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; bypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            lung surfactant in a subject's tissue, or treating bronchoconstriction, lung inflammation, lung allergies, or a respiratory disease or condition. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel pharmaceutical composition, which first active agent comprising an oligonucleotide antisense to the intelation codon, coding region, 5' or 3' end genomic flanking region, 5' or 7' end genomic flanking region, 5' or regions within 2-10 nucleotide, or regions within 2-10 nucleotide.
Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABZ90836;
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                                                                                                 WPI; 2003-229219/22
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Tang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 A; 0 C; 2 G; 0 T; 0 U; 0 Other;
                                                                                                                                                           Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.8; DB 1;
Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  disease; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Migmatches
                                                                                                                                                           Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 20;
                                                                                                                                                                                               Pabalan
                                                                                                                                                                                            Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                         Aguilar D;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         regions,
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corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone

SEQ ID NO 6078; 872pp; English

The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cycostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC for enhancing the prophylactic or therapautic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of or reducing sensitivity to adenosine, reducing levels of adenosine creceptor, producing bronchodilation, increasing levels of ubiquinone or CC lung inflammation, lung allergies, or a respiratory disease or condition.

CC Note: The sequence data for this patent is not represented in the printed as pecification, but was obtained in electronic format directly from WIPO as termination. at ftp.wipo.int/pub/published_pct_sequences

Sequence 20 BP; 6 A; 2 C; 9 G; 3 T; 0 U; 0 Other;

Matches Query Match Best Local (1 Similarity Conservative 0.3%; 0; Mismatches Score 15.8; DB 1; Pred. No. 7.7e+02; 0; Mismatches 2; Length 20 0, Gaps 0

밁 δ 2945 GAAACCTGAAGGAGCTGGA 2963 1 GAAAGCTGAAGGTGCTGGA 19

ABZ85670 standard; DNA; 20 ₽P

ABZ85670

17-OCT-2003 (first entry)

Human oligonucleotide sequence.

RESULT 564
ABZ85670/C
ID ABZ8566
XX ABZ856
XX ABZ856
XX Human
XX Homo 8
XX 1020
ZX Homo 8
XX Homo 8
XX Homo 8
XX Homo 9
ZX Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.

Homo sapiens

WO200285308-A2

31-OCT-2002

23-APR-2002; 2002WO-US013135

24-APR-2001; 2001US-0286137P

(EPIG-) EPIGENESIS PHARM INC.

ŝ Li Y, Tang Ļ Sandrasagra A, , Shahabuddin Katz E, S; Pabalan Ç Aguilar

WPI; 2003-229219/22.

Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its

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RESULT 565
ABZ87765
     5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel pharmaceutical composition, which has a CC first active agent comprising an oligonucleotide antisense to the CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC or enhancing the prophylactic or therapeutic respiratory effect of an antiinflammatory steroid in a subject, for reducing or depleting levels of, or reducing sensitivity to adenosine, reducing levels of adenosine conceptor, producing bronchodilation, increasing levels of ubiquinone or CC lung surfactant in a subject's tissue, or treating bronchoconstriction, unterinted specification, but was obtained in electronic format directly from WIPO care for wiso intronucled in the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                            Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; bypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its
                                                       WPI; 2003-229219/22
                                                                                       Miller S,
                                                                                                                                                                                                                                                                                        WO200285308-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human oligonucleotide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ87765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABZ87765 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              corresponding RNAs, ubiquinone.
                                                                                                                                                                                24-APR-2001; 2001US-0286137P
                                                                                                                                                                                                                  23-APR-2002; 2002WO-US013135
                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5403 AAAAAAGAAAAATGAAAA 5421
                                                                                                                                                                                                                                                                                                                                                             inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19
                                                                                                                                             EPIGENESIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAAATAGAAAAAAAAGAAAA 1
                                                                                       Li Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 1 A; 2 C; 0 G; 17 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                         Sandrasagra A,
L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             912; 872pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.8; DB 1; Length 20; Pred. No. 7.7e+02;
                                                                                       Katz E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                           Pabalan
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                                                                                                        Aguilar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                        Disclosure; SEQ ID NO 3007; 872pp; English
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The invention relates to a novel pharmaceutical composition, which has a CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC initiation codon, coding region, 5' or 3' end genomic flanking regions, CC 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of CC junctions of genes encoding a polypeptide associated with lung and/or CC nasal airway dysfunction and a second active agent comprising an CC antiinflammatory steroid and ubiquinone. A composition of the invention CC immunosuppressive, and cytostatic activity. The composition may have a CC use in antisense gene therapy. The composition is useful for treating or preventing a respiratory, lung or malignant disease or condition, also CC or enhancing the prophylactic or therapeutic respiratory effect of an CC entiinflammatory steroid in a subject, for reducing levels of adenosine cof or reducing sensitivity to adenosine, reducing levels of adenosine cor lung inflammation, lung allergies, or a respiratory disease or condition. CC Note: The sequence data for this patent is not represented in the printed contribution, but was obtained in electronic format directly from WIPO CC at fer, wino, intronbhodial active agentatory. at ftp.wipo.int/pub/published_pct_sequences

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                                                               RESULT 566
                                                                                                                                                 Query Match
Best Local
                                                                                                                                        Matches
                     ABZ98968
 17-OCT-2003
                                          ABZ98968 standard; DNA; 20
                                                                                                                  2833 GAGGGCAGGCACAGAT 2851
                                                                                                                                                 Similarity
                                                                                               GAGGGCACGGAGCACAGAT 20
                                                                                                                                       0.3%;
ilarity 89.5%;
Conservative
(first entry)
                                                                                                                                       0;
                                                                                                                                                 Score 15.8;
Pred. No. 7.
                                                                                                                                         Mismatches
                                                                                                                                                .7e+02;
                                                                                                                                                           DB 1; Length 20;
                                                                                                                                         Indels
                                                                                                                                        ,,
                                                                                                                                        Gaps
                                                                                                                                         0
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Sequence 20 BP; 7 A; 4 C; 8 G; 1 T; 0 U; 0 Other;

ABZ98968

ID ABZ;

XX

AC ABZ

XX

DT 17-
XX

Hum

XX

HW

KW ant

KW ant

KW ant

KW ant

KW ant

XX

PN WO

XX

PN WO

XX

PN WO

XX

PN WO

XX

PF 23

XX

PR 24

XX

PN WO

XX

PN PI W

XX

PN W

XX

PN PI W Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; immunosuppressive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; WO200285308-A2 Homo sapiens. inflammation; respiratory disease; ds.

Human PDE4A oligonucleotide sequence.

24-APR-2001; 2001US-0286137P 23-APR-2002; 2002WO-US013135

WPI; 2003-229219/22. (EPIG-) EPIGENESIS PHARM INC Li Y, Tang F, Sandrasagra A, L, Shahabuddin Katz E, Pabalan J,

Pharmaceutical composition respiration, has oligo(s) a

n for treating ailments associated with impaired antisense to specific gene(s) or its

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RESULT 567
ACC47647/c
ID ACC47647/c
XX ACC476
XX ACC476
XX ACC476
XX Human
XX Human
XX Human
XX Chromo
KW Chromo
KW Chromo
KW Duchel
KW Duches
KW 
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; insulin-like growth factor binding protein 5; IGFBP5; IBP5; chromosome 2q33-34; IGF signal transduction; IGF regulation; apoptosis; bone growth stimulator; hyperproliferative disorder; cancer; tumour; breast; prostate; pancreas; neuroendocrine; inflammatory disorder; colitis, developmental disorder; growth disorder; disorder; buchenne muscular dystrophy; metabolic disorder; diabetes; osteoporosis; cytostatic; antiinflammatory; expression inhibition; phosphorothioate; antisense oligonucleotide; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACC47647 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or ubiquinone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human IGFBP5 phosphorothioate antisense oligonucleotide, SEQ ID NO:23.
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                                      07-OCT-2002; 2002WO-US032060
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-SEP-2003
                                                                                                                                                                                                               WO2003030826-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3215 GACTGCAGCTGGTCAGCTG 3233
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GACTGCAGCACGTCAGCTG 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                           /mod_base= OTHER
/mod_base= OTHER
/note= "This oligonucleotide has a phosphorothioate
/note= "This oligonucleotide has a phosphorothioate
backbone and 2 'methyoxyethyl (2'-MOE) wings at the
and 3' ends, which are 5 nucleotides in length. Also
corrections residues are 5-methyloytosines
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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Pred. No. 7.7e+02;
0; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 20
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                                                                                                                                                                                                                                                                                                                                                                                    wings at the 5'
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                                                                                                                                                                                                                                                                                                                                    Also all
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modified_base

/*tag=

. 20

Location/Qualifiers

Mus musculus

disease;

88;

mouse

antisense; p38 mitogen activated protein kinase; p38 MAPK; inflammatory disease; autoimmune disease; rheumatoid arthritis;

03-JUN-2004 ADM34276

(first entry)

Моиве р38

MAPK antisense oligonucleotide #3.

ADM34276 standard; DNA; 20

ВP

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ADM34276
IID ADM1
XX
AC ADM1
XX
ON 03-0
XX
ON Moun
XX
KW Annt
XW Annt
XW hear
XX
OS Mus
XX
FH Key
FT modd
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                                                                                                                                                                                                                                                                                                                                                                                 RESULT 568
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CC oligonucleoxides targeted to the human insulin-like growth factor binding conteins 5 (IGFBP5) gene, which inhibit its expression. The antisense colligonucleoxides were designed to target different regions of human colligonucleoxides were designed to target different regions of human colligonucleoxides were designed to target different regions of human colligonucleoxides were analysed for their effect on IGFBP5 expression by component of their effect on IGFBP5 expression by component of the IGF system in bone, having a high college of the insulin-like growth factor superfamily, which are involved in the college of the insulin-like growth factor superfamily, which also mediate IGF-college of the IGF system in bone, having a high component of the IGF system in bone, having a high contains, and appears to act as a growth factor, stimulating bone content rissues, such as kidney, liver, gut endothelium, lung tubules and contert tissues, such as kidney, liver, gut endothelium, lung tubules and contertyme, meninges, notochord, muscle and tongue. Its levels are also increased in inflamed colon smooth muscle and tongue. Its levels are also concreased in inflamed colon smooth muscle calls in an experimental model concreased in inflamed colon smooth muscle calls in an experimental model concreased in inflamed colon smooth muscle calls upon treatment concretion in the strongens used to abolish tamoxifen resistant proliferation. The oligonucleoxides to abolish tamoxifen resistant proliferation. The oligonucleoxides of the invention are useful for diagnosis, colitis, developmental or growth disorders, such as growth as to contert of the breast, colitis, developmental or growth disorders (e.g., Duchenne contents), or metabolic disorders (e.g., diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antisense oligonucleotides for modulating insulin-like growth factor binding protein 5 gene expression, useful for preventing or treating cancers, inflammatory disorders, developmental disorders or metabolic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequences ACC47637-ACC47667 represent phosphorothioate antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; Page 76; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-OCT-2001; 2001US-00975123.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1634 AGCTGGCCCAGTCCAAGGT 1652
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17; Conserv
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nilarity 89.5%;
Conservative
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Pred. No. 7.7e+02;
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RESULT 569
ABD24268/c
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DE Human
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                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                  The invention relates to an antisense oligonucleotide 8-30 nucleobases in length targeted to the 5'-untranslated region, translational start site, translational termination region or 3'-untranslated region of a nucleic acid molecule encoding a p38 mitogen activated protein kinase (MAPK), where the antisense compound inhibits the expression of the p38 MAPK. The antisense oligonucleotide is useful for inhibiting the expression of p38 MAPK in cells or tissues. It is also useful for treating an animal having a disease or condition associated with p38 MAPK, e.g. an inflammatory or an autoimmune disease (e.g. rheumatoid arthritis) or a heart disease. In addition, the compound is used for diagnostics, prophylaxis, or as research reagents or kits. The present sequence represents a p38 MAPK antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotides for modulating p38 mitogen activated protein kinase (MAPK) expression, useful for diagnosing, preventing or treating diseases associated with p38 MAPK, e.g. inflammation or heart
 Human calmodulin 2-derived oligonucleotide SEQ ID
                             29-JUL-2004
                                                         ABD24268
                                                                                  ABD24268 standard; DNA; 20
                                                                                                                                                                                                                                                           Sequence 20
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15-AUG-2000; 2000US-00640101.
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NERO P.
MCKAY R.
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/note= "2'-methoxyethoxy
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Pred. No. 7.7e+02
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Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antinflammatory; antisethmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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Homo sapiens.

WO200285309-A2.

31-OCT-2002.

23-APR-2002; 2002WO-US013143.

24-APR-2001; 2001US-0286036P

(EPIG-) EPIGENESIS PHARM INC.

Nyce JW, Li Y, Sandrasagra A, Katz E, Pabalan J, Aguilar D; Miller S, Tang L, Shahabuddin S;

WPI; 2003-093058/08.

pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and bronchodilating agent.

Claim 15; SEQ ID NO 3280; 763pp; English.

CC comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating conchoconstriction, respiratory tract inflammation, allergies and concentrated depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating conjuncted tides are derived from a gene encoding or regulating construction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery construction also describes a kit, that comprises: (a) a delivery construction in separate containers, (b) the oligonucleotides, (c) constructions for adding a carrier and for use of the kit. The composition cof the invention has antiallergic, antialinflammatory, antiasthmatic, analysis, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition, allergies and/or bronchoconstriction and/or lung inflammation, allergies, asthma, impeded respiration, respiratory conflammation, allergies, asthma, impeded respiration, respiratory conflammation, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system of the oligonucleotides into products that free adenosine into the system of the content of the oligonucleotides into the system of the content

Sequence 20 BP; 3 A; 7 C; 9 G; 1 T; 0 U; 0 Other;

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Query Match
0.3%; Score 15.8; DB 1; Length 20;
Best Local Similarity 89.5%; Pred. No. 7.7e+02;
Matches 17; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
Oy 2641 CTGCAGCTGCTGCAGC 2659
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24-APR-2001; 2001US-0286036P 23-APR-2002; 2002WO-US013143

Miller S, Li Y, Tang Sandrasagra A, Shahabuddin 5 Katz 'n Pabalan ç

'n

(EPIG-)

EPIGENESIS

PHARM INC

WPI; 2003-093058/08.

Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targ nucleic acids associated with lung airway or lung dysfunction bronchodilating agent. or lung dysfunction, targeted to

Claim 15, SEQ ID NO 6078; 763pp; English.

CC comprising oligonucleotides, effective for alleviating CC pronchoconstriction, respiratory tract inflammation, allergies and CC reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, CC surfactant depletion or hyposecretion, when administered to a mammal. The CC oligonucleotides are derived from a gene encoding or regulating CC expression of a target polypeptide associated with lung airway or lung CC dysfunction or cancer and can be anti-sense to the corresponding mRNA. CC The invention also describes a kit, that comprises: (a) a delivery CC device, in separate containers, (b) the oligonucleotides, (c) constructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The CC inflammation, allergies and/or bronchoconstriction and/or lung conflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction are associated conflammation, allergies, asthma, impeded respiration, respiratory thyperrension, emphysema, chronic obstructive pulmonary disease, pulmonary disease, pulmonary conditions, bronchitis or cancer.

CC The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of thymidines present in the target RNA serves to prevent the breakdown of invention describes a novel composition rising oligonucleotides, effective for al <u>a</u> ρ first active agent,

This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, surfactant depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung aliway or lung dysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a kit, that comprises: (a) a delivery device, in separate containers, (b) the oligonucleotides, (c) instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic,

analgesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition is useful for preventing or treating a respiratory, lung or malignant disease. The administered

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the oligonucleotides into products that free adenosine into the system e.g., lung, brain, heart, kidney, etc, tissue environment and thereby, prevent any unwanted effects due to it
                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, an
                                                                                                                                                                                          Claim 15; SEQ ID NO 912; 763pp; English.
                                                                                                                                                                                                                 bronchodilating agent.
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L, Shahabuddin
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ARESULT 57
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AD ARBOLT 644
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AC ARBOLT 
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                                                                                                                                                                                                                                        Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABD21764 standard; DNA; 20
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L, Shahabuddin
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This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, allergies and

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Shahabuddin

Claim

15;

SEQ ID NO 776;

763pp; English

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                                 Nyce JW,
Miller S,
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                                                                                                                                   EPIGENESIS PHARM INC
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                                 Li Y,
Tang
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                                                                   Sandrasagra A,
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Pred. No. 7.7e+02;
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                                                                   Katz E,
                                                                      Pabalan J,
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                                                                                                                                                                                                                                                                         Human calmodulin 2-derived oligonucleotide SEQ ID 3007.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the oligonucleotides into products that free adenosine into e.g., lung, brain, heart, kidney, etc, tissue environment ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 15; SEQ ID NO 14210; 763pp; English
                                                                                 pulmonary transplantation
                                                                                                                    beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprising oligonucleotides, effective for alleviating
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                                                                                                      emphysema; chronic obstructive pulmonary disease;
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17; Conserv
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                                                                                 rejection; ss;
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(A) or (A) receptors,
Tammal. The
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WO200285309-A2

Homo sapiens.

RESULT 575
ADF66213/c
ID ADF662
XX
AC ADF662
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DT 26-FEB
XX
DE Ian5 9
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ADF66213

ADF66213 standard; DNA; 20

ВP

Ian5 gene related PCR primer SEQ ID NO:32

26-FEB-2004 (first entry)

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N

2833 GAGGGCAGCAGCAGAT 2851

0; Mismatches

Indels

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Gaps

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CC dysfunction or cancer and can be anti-sense to the corresponding mRNA.

CC The invention also describes a kit, that comprises: (a) a delivery

CC device, in separate containers, (b) the oligonucleotides, (c)

CC instructions for adding a carrier and for use of the kit. The composition

CC of the invention has antiallergic, antiinflammatcory, antiasthmatic,

CC analgesic, hypotensive, immunosuppressive and cytostatic activity, is a

CC treating a respiratory, lung or malignant disease. The administered

CC composition comprises oligo and is administered to reduce the production

CC or availability, or to increase the degradation of the target mRNA or to

CC reduce the amount of target polypeptide present in the lungs. The

CC conflammation, allergies and/or bronchoconstriction and/or lung

CC inflammation, allergies and/or bronchoconstriction and/or lung

CC inflammation, allergies, asthma, impeded respiration, respiratory

CC distress syndrome, pain, cystic fibrosis, allergic rhinitis, pulmonary

CC transplantation rejection, pulmonary infections, bronchitis or cancer.

CC thymothemist present in the target RNA serves to prevent the breakdown of

CC the oligonucleotides into products that free adenosine into the system

CC prevent any unwanted effects due to it
  Best Local Similarity Matches 17; Conserv
                    Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              surfactant depletion or hyposecretion, when administered to a mammal. oligonucleotides are derived from a gene encoding or regulating expression of a target polypeptide associated with lung airway or lung
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprising oligonucleotides, effective for alleviating bronchoconstriction, respiratory tract inflammation, aller reducing adenosine sensitivity, levels of adenosine (A) or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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                                                                              Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    bronchodilating agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-APR-2002; 2002WO-US013143
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Li Y, Sar
, Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 3007; 763pp; English
0.3%;
ilarity 89.5%;
Conservative
                                                                              BP; 7 A; 4 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        describes a novel composition (a) a first active agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sandrasagra A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shahabuddin
                      Score 15.8; DB 1
Pred. No. 7.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Katz E,
                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pabalan J,
                                       Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 allergies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Aguilar D;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptors,
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                                                                                                                                                                     The present invention describes a GAD65 polypeptide comprising an E517P cmutation. The polypeptide is characterised by decreased specific binding to an antibody selected from GAD6, MICA-1, MICA-3, MICA-4 and MICA-6, CC where the decreased binding is relative to a corresponding GAD65 cC polypeptide not having the E517P mutation. Also described: (1) methods CC encode the rat Ian5(4) polypeptide; or or risk of type 1 diabetes in a subject; CC (2) an isolated mucleic acid selected from: (a) mucleic acids which encode the cart Ian5(1yp) polypeptide; or (c) full length complements of the nucleic acids of (a) or (b); (3) an antibody that: (a) specifically binds to rat Ian5(1yp) polypeptide; or (c) full length complements of the nucleic acids of (a) or (b); (3) an antibody is not immunologically cross-reactive with human Ian5 or mouse Ian5 polypeptide; or (b) specifically cross-reactive with human Ian5 or mouse Ian5 polypeptide; or (b) specifically cross-reactive with numan Ian5 or mouse Ian5 polypeptide; or mouse Ian5 for mouse Ian5
밁
                                                                                Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New GAD65 polypeptide comprising an E517P mutation, useful for detecting the presence of or risk of developing type 1 diabetes or a related disorder in human subjects.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAD65; type 1 diabetes; rat; Ian5(+); Ian5(lyp); Ian5; antidiabetic;
gene therapy; diabetes; PCR primer; ss.
                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-MAY-2003; 2003WO-US017206
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO2003102147-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; SEQ ID NO 32; 87pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-053464/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UNIW ) UNIV WASHINGTON
                                       572 AGAAGGAGGAGCTGAAGGA 590
19 AGAAGAGGGAGCTGAAGGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gp.
                                                                                                      Similarity
                                                                                                                                                                 BP; 1 A; 9 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ď,
                                                                                                    0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Macmurray
                                                                                0;
                                                                                                                         Score 15.8;
                                                                                                      Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P
                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ettinger RA,
                                                                                                      7.7e+02
                                                                                                                         DB 1; Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Moralejo D;
                                                                                0
                                                                                Gaps
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RESULT 576 ADG75970/c

RESULT 577
ADH26716/c
ID ADH267
XX
AC ADH267
XX
DT 11-MAR

ADH26716 standard; DNA; 20

ADH26716;

11-MAR-2004

(first entry)

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                             S
                                                                                                                                                 CC contain a non-palindromic sequence motif. Specifically, it refers to DNA CC oligonucleotides (without a CpG motif), which can stimulate an immune CC response in animals of the order of primate, including humans. The immune CC response is characterised by the proliferation, differentiation, cytokine ccluding humans of the order of primate, including humans. The immune CC and antibody production in B-cells, as well as cell differentiation and CC cytokine production in plasmacytoid dendritic cells. The present CC invention describes immunomodulator compositions that also comprise an CC antigen selected from, for example, viruses, bacteria, parasites, tumour CC cells and glycolipids. As such, these DNA oligos can be used in gene CC therapy for inducing B-cell activation, treating, preventing or CC ameliorating an immune system disorder or a tumoural disease including CC chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell CC carcinoma. This oligonucleotide sequence is an immunostimulatory CC phosphorothioate non-CpG variant DNA oligo, used to determine the effect CC of oligo size on B cell proliferation and IL6 secretion in an CCC exemplification of the invention.
                                                               Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      proliferation; differentiation; cytokine; antibody production; plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma; renal cell carcinoma.
                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30-MAY-2003; 2003WO-EP005691.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to novel immunostimulatory oligonuclectides that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 5; SEQ ID NO 72; 139pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-053333/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30-MAY-2002; 2002CA-02388049.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003101375-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADG75970 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lopez RA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (IMMU-) IMMUNOTECH SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-DEC-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Immunostimulatory non-CpG phosphorothioate DNA oligo IMT189 SeqID72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11-MAR-2004
                                5403 AAAAAAGAAAAATGAAAA 5421
                                                                                                                                                 BP; 1 A; 1 C; 1 G; 17 T; 0 U; 0 Other;
                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry
                                                                                         0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP
                                                                       0,
                                                                                         Score 15.8;
Pred. No. 7.
                                                                           Mismatches
                                                                                       .7e+02;
                                                                                                                B
                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      immune response;
                                                                           Indels
                                                                                                                20
                                                                           0,
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                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antisense oligonuclectides inhibiting the expression of phosphoinositide-3-kinase, regulatory subunit 4, p150, useful for preventing or treating diseases associated with the subunit, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PI3 kinase; ss; antisense oligonucleotide; phosphorothioate li
2'-O-methoxyethyl sugar moiety: free thylcytosine;
hyperproliferative disorder; cancer; Chediak-Higashi syndrome;
neurodegenerative disorder; metabolic disorders; inflammation;
cytostatic; immunomodulator; neurodegenerative; antimicrobial;
Human; phosphoinositide-3-kinase regulatory subunit 4; p150; PI3K; PI3 kinase; ds; antisense oligonucleotide; phosphorothioate linkage;
                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 15; SEQ ID NO 51; 62pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       preventing or treating diseases associated hyperproliferative disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-051923/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-MAY-2002; 2002US-00160786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-MAY-2002; 2002US-00160786.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2003225013-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human PI3K regulatory subunit 4,
                                                                                                                                                                 ADH26784 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antiinflammatory.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                            376 GGTGCCCTGGGATTTATAA 394
                                                      PI3K
                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     phosphoinositide-3-kinase regulatory subunit 4; p150; PI3K;
                                                                                                                                                                                                                                                          GGTGCCGTGGGATTTATCA 2
                                                      regulatory subunit 4,
                                                                                                                                                                                                                                                                                                                                                                                                      BP, 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dobie KW;
                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                      A; 7 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                 0.3%;
89.5%;
                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                        Score 15.8; Db .,
                                                                                                                                                                                                                                                                                                                                Mismatches
                                                      p150 DNA target region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            p150 DNA antisense oligonucleotide #41.
                                                                                                                                                                                                                                                                                                                                                                  DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                0,
                                                                                                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                      RESULT 579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a compound targeted to a nucleic acid molecule encoding human phosphoinositide-3-kinase (PI3K) regulatory subunit 4, p150. The compound is an antisense oligonucleotide that specifically hybridises with a nucleic acid molecule encoding PI3K regulatory subunit 4, p150 and inhibits expression of the polypeptide. The antisense oligonucleotide comprises at least one modified internucleoside linkage oligonucleotide comprises at least one modified sugar molety, preferably a 2'-0-methoxyethyl sugar molety, or at least one modified nucleobase comprising a 5-methylcytosine. The antisense compounds are useful for modulating the expression of PI3K regulatory subunit 4, p150 and for preventing or treating hyperproliferative disorders (i.e. cancer), Chediak-Higashi syndrome, neurodegenerative disorders and metabolic disorders. These may also be used in research and diagnostics and in preventing or delaying infection or inflammation. This sequence represents a PI3K regulatory subunit 4, p150 DNA antisense
                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotides inhibiting the expression of phosphoinositide-3-kinase, regulatory subunit 4, p150, useful preventing or treating diseases associated with the subunit, e hyperproliferative disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2'-O-methoxyethyl sugar moiety; 5-methylcytosine;
hyperproliferative disorder; cancer; Chediak-Higashi syndrome;
neurodegenerative disorder; metabolic disorders; inflammation;
cytostatic; immunomodulator; neurodegenerative; antimicrobial;
                                 inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's phosphorothioate backbone; 2'-methoxyethyl; 2'-MOB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAY-2002; 2002US-00160786.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                antisense oligonucleotide; glucocorticoid receptor; infection;
                                                                                                                 Human glucocorticoid receptor-specific antisense oligonucleotide
                                                                                                                                                      25-MAR-2004
                                                                                                                                                                                                                      ADH65486 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 119; 62pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-MAY-2002; 2002US-00160786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antiinflammatory.
                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2004-051923/05
                                                                                                                                                                                                                                                                                                                                        376 GGTGCCCTGGGATTTATAA 394
                                                                                                                                                                                                                                                                                                       1 GGTGCCGTGGGATTTATCA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MS,
                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A; 4 C; 7 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                    (first entry)
                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
                                                                    formation;
                                                                                                                                                                                                                      20
                                                                                                                                                                                                                      ВÞ
                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                       Score 15.8; DB 1;
Pred. No. 7.7e+02;
                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                    syndrome; human;
                                                                                                                                                                                                                                                                                                                                                                           0
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                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Homo sapiens

88

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RESULT 580
ADH6577/C
ID ADH657
AC ADH657
AC ADH657
AC ADH657
XX

DT 25-MAR
XX

ANTIBE
KW INFLAM
CARDIO
KW EARDIO
EARDI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   밁
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTS: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE.
                                 WPI; 2004-035034/03
                                                                                       Crosby SD,
                                                                                                                                                                                                                                                                                                                                                                                 WO2003099215-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human glucocorticoid receptor-specific antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADH65757;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADH65757 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention comprises an antisense oligonucleotides that are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17;
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                                                                                                                                                PHARMACIA CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                       Nalseth AE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP; 3 A; 12 C; 0 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                         2002US-0381857P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
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Pred.
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No. 7
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New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                         Claim 4; SEQ ID NO 2591; 985pp; English.
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to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumonur formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothicate backbone. The invention comprises an antisense oligonucleotides that are targeted

Sequence 20 BP; 2 A; 11 C; 0 G; 7 T; 0 U; 0 Other;

Query Match Best Local S Matches 17 Similarity Conservative 0.3%; <u>,</u> Score Pred. Mismatches No. 7 .7e+02 DB 1; Length Indels 20; 0; Gaps 0;

밁 S 2564 AGGGGGAGAGAGAGATGGA 2582 AGGGGAAGAGGGAGATGGA 1

ADH66756 standard; DNA;

20

ADH66756;

(first

25-MAR-2004

entry)

Human glucocorticoid receptor-specific antisense oligonucleotide #3590.

cardiovascular disorder; hyperlipidaemia; Cushing's syndrome; human; phosphorothioate backbone; 2'-methoxyethyl; 2'-MOE. antisense oligonucleotide; glucocorticoid receptor; inflammation; tumour formation; diabetes; infection;

sapiens.

WO2003099215-A2

04-DEC-2003

20-MAY-2003; 2003WO-US016084.

20-MAY-2002; 2002US-0381857P.

(PHAA) PHARMACIA CORP

SD, Nalseth

2004-035034/03.

New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, ob cardiovascular disorder, hyperlipidemia or Cushing's syndrome. obesity,

Claim 4 SEQ ID NO 3590; 985pp; English.

RESULT 581
ADH65756/C
ID ADH6577
XX ADH6577
AC ADH6677
XX 25-WAR
XX 25-WAR
XX antise
KW inflam
KW cardic
KW phosph
XX Homo s
XX Homo s
XX 20-WAN
XX 20-WAN
XX 20-WAN
XX 20-WAN
XX Crosb
YN WPI; i
XX New ar
PT mammal
PT cardic
XX New ar
PT mammal
XX New ar
PT mammal
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CC to nuc
CC antise
CC coligor
CC collagy
CC co The invention comprises an antisense oligonucleotides that are targeted to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothioate backbone.

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RESULT 583
ADJ46820/c
ID ADJ468
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Best Local Similarity
Matches 17; Conserva
                                                                                                                                                                                       RESULT 582
ADH67178
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Best Local S
Matches 17
                                                                                                                                                                                                               The invention comprises an antisense oligonucleotides that are targeted to nucleic acids encoding a mammalian glucocorticoid receptor. The antisense oligonucleotides of the invention are useful for preventing or delaying infection, inflammation or tumour formation. The antisense oligonucleotides are also useful for treating diabetes, obesity, cardiovascular disorders, hyperlipidaemia or Cushing's syndrome. The present DNA sequence represents an antisense oligonucleotide that targets the human glucocorticoid receptor gene. NOTE: The present sequence contains 2'-methoxyethyl (2'-MOE) wings and a phosphorothicate backbone.
                                                                                                                                                                                                                                                                                                                                                                                             New antisense compound targeted to a nucleic acid molecule encoding mammalian glucocorticoid receptor, useful for treating diabetes, obesity, cardiovascular disorder, hyperlipidemia or Cushing's syndrome.
                                                                                                                                                                                       Sequence 20 BP; 3 A; 11 C; 0 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense oligonucleotide; glucocorticoid receptor; infection; inflammation; tumour formation; diabetes; obesity; cardiovascular disorder; hyperlipidaemis; Cushing's syndrome; human; ss; phosphorothioate backbone; 2'.methoxyethyl; 2'.MOE.
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les 17; Conserv
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                                                                                                  217 CACCACATCTCCCCTCACC 235
                                                                                                                                                                                                                                                                                                                                                                   4; SEQ ID
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                                                                        CATCACATCTCCCCTCTCC 19
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nilarity 89.5%;
Conservative
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Pred. No. 7.7e+02.
0; Mismatches
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Pred. No. 7.7e+02.
0; Mismatches
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ADJ46820 standard; DNA; 20

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05-FEB-2004

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RESULT 584
ADJ467474
ID - ADJ467
XX ADJ467
XX O6-MAY
XX Human
XX human;
XW Homo 8
OS Synthe
XX Synthe
XX US2004
XX XX
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New compound, particularly an antisense oligonucleotide targeted to a nucleic acid encoding KIAA1531, useful for treating cancer, chronic inflammation or conditions involving hyperactivation of angiogenesis.
                                             US2004023378-A1
                                                                                                                  human; KIAA1531; hyperproliferative disorder;; cancer; angiogenesis hyperactivation; chronic inflammation; ss
                                                                                                                                                              Human
                                                                                                                                                                                           06-MAY-2004
                                                                                                                                                                                                                                                  ADJ46745 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 6 A; 1 C; 9 G; 4 T; 0 U; 0 Other;
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                                                                         Synthetic
                                                                                        Homo sapiens
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                                                                                                                                                            KIAA1531 antisense oligonucleotide ISIS #208169.
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                                                                                                                                                                                           (first entry)
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Pred. No. 7.
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Matches
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                                                                                                                                                                                                                                                                                                   interleukin; IL-4 receptor; IL-5 receptor; lung disease; airway inflammation; allergy; asthma; impeded respiration; cystic fibrosis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New compound, particularly an antisense oligonucleotide targeted to a nucleic acid encoding KIAA1531, useful for treating cancer, chronic inflammation or conditions involving hyperactivation of angiogenesis.
Novel single or multiple target oligonucleotide anti-sense to e.g. initiation codons and introns of respiratory disease-relevant genes e.g., CCR1, RANTES, MCP4, useful for prophylaxis or treating respiratory
                                                                                Nyce JW, Tang
Shahabuddin S,
                                                                                                                                                                                                                                   WO2004011613-A2
                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide associated to PDE4A #134.
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H, Cong H;
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                                                                                              Aguilar D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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RESULT 586
ADJ18373/C
ID ADJ18373/C
ID ADJ183
XX ADJ183

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (COPD), allergic rhinitis (AR), acute respiratory distress syndrome (ARDS), pulmonary hypertension, lung inflammation, bronchitis, airway obstruction. The present sequence represents an oligonucleotide of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease e.g., asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human; ss; liver related homologue-1; LRH1; NR5A2; antisense; phosphorothioate; 2' MOE; breast cancer; dyelipidaemia; atherosclerosis; low HDL, high density lipoprotein; high LDL, hypercholesterolaemia; gall stone; triglyceridaemia; obesity; hepatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADJ18373 standard;
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                               08-JAN-2004.
                                                                                            WO2004003201-A2
                                                                                                                                                                                                                                                                          modified_base
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modified_base
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Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antiinflammatory;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    antiarteriosclerotic; anorectic; hepatotropic; litholytic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hepatocellular carcinoma; aromatase; cytostatic; antilipaemic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3215 GACTGCAGCTGGTCAGCTG 3233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GACTGCAGCACGTCAGCTG 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA oligo used to modulate human LRH1 expression SeqID 2923.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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/mod_base= OTHER
^murR= p
                                                                                                                                                                                                                                                                                          /mod_base= OTHER
/note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides. All
/vridine nucleobases are 5-methylcytidine."
                                                                                                                                                  /note= "OTHER= 2' methoxyethyl (2' MOE) nucleotides.
rytidine nucleobases are 5-methylcytidine."
                                                                                                                                                        cytidine
                                                                                                                                                                                                                 /*tag=
/mod_ba
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 virucidal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.3%; Score 15.8; DB 1;
89.5%; Pred. No. 7.7e+02;
                                                                                                                                                                                                                 base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER= phosphorothicate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 20;
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RESULT 587
ADJ18838/c
ID ADJ188
XX ADJ188
XX ADJ188
XX Antise
XX Antise
XX Antise
XX Homan;
KW human;
KW possph
KW low HD
KW low HD
KW antisx
KW hepatc
KW antisx
KW Synthe
XX
FH Key
FT modifi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to novel antisense compounds useful for modulating CC the expression of liver related homologue-1 (LRH1) and splice variants CC thereof. Specifically, it refers to compositions 93 0 nucleobases in CC length that target a portion of an active site on the nucleic acid combecule encoding LRH1 (also known as NR5A2). LRH1 is a monomeric orphan cc nuclear receptor protein that functions as a tissue specific transcription factor. The present invention describes antisense coligonucleotides that comprise at least one modified internucleoside cc linkage, a phosphorothicate linkage; at least one modified sugar moiety, ca 2'-O-methoxyethyl (2' MOS) and at least one modified nucleobase, a 5-cc methylcytidine. These antisense compounds are useful for treating or CC diagnosing a disease associated with LRH1, such as breast cancer, cc dyslipidaemia, atherosclerosis, low HDL (high density lipoprotein), high criditis, as well as hepatocoellular carcinoma or a condition associated with aromatase activity. Accordingly, these compositions exhibit controlic, antilinfammatory and virucidal activities. This cc oligonucleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRH1 protein of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                          human; ss; liver related homologue-1; LRH1; NR5A2; antisense; phosphorothioate; 2' MOB; breast cancer; dyslipidaemia; atherosclerosis; low HDL; high density lipoprotein; high LDL; hypercholesterolaemia; gall stone; triglyceridaemia; obesity; hepatitis; hepaticis; antilipaemic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New antisense oligonucleotides targeted to a nucleic acid encoding related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                           Synthetic.
                                                                                                                                      antiarteriosclerotic; anorectic;
antiinflammatory; virucidal.
                                                                                                                                                                                                                                                                                                                                     20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 15; SEQ ID NO 2923; 909pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-JUL-2003;
                  modified_base
                                                                                                                                                                                                                                                                                               Antisense DNA oligo used to modulate human LRH1 expression SeqID
                                                                                                                                                                                                                                                                                                                                                                                                                 ADJ18838 standard; DNA; 20 BP
                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGAATGTCTTCTAATTCAG 2
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Location/Qualifiers
1. .20
/*tag= b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.8; DB 1; Pred. No. 7.7e+02;
                                                                                                                                                        hepatotropic; litholytic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length
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RESULT 588
ADK74191/c
ID ADK741
XX
AC ADK741
XX
AC ADK741
XX
DT 20-MAY

ADK74191 standard;

DNA; 20

ВÞ

20-MAY-2004 ADK74191;

(first entry)

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This invention relates to novel antisense compounds useful for modulating CC the expression of liver related homologue-1 (IRH1) and splice variants CC thereof. Specifically, it refers to compositions 8-30 nucleobases in CC length that target a portion of an active site on the nucleic acid CC molecule encoding IRH1 (also known as NR5A2). IRH1 is a monomeric orphan CC nuclear receptor protein that functions as a tissue specific CC transcription factor. The present invention describes antisense CC oligonucleotides that comprise at least one modified internucleoside CC linkage, a phosphorothicate linkage; at least one modified sugar moiety, CC a 2'-O-methoxyethyl (2' MOB) and at least one modified nucleobase, a 5-CC methylcytidine. These antisense compounds are useful for treating or CC diagnosing a disease associated with IRH1, such as breast cancer, CC dyslipidaemia, atherosclerosis, low HDL (high density lipoprotein), high CLDL (low density lipoprotein), hypercholesterolaemia, gall stones, CC triglyceridaemia, obesity, hepatitis B virus-mediated acute or chronic hepatitis, as well as hepatocellular carcinoma or a condition associated with the compositions are such that contains a such that contains a such that contains a such that the compositions are such that contains a such that composition as such that composition and contains a such that composition as such that compos
                                                                                                                                                  Matches
                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                with aromatase activity. Accordingly, these compositions exhibit cytostatic, antilipaemic, antiarteriosclerotic, anorectic, hepatlitholytic, antiinflammatory and virucidal activities. This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense oligonucleotides targeted to a nucleic acid encoding related homologue-1 (LRH1), useful for treating breast cancer, dyslipidemia, atherosclerosis, hypercholesterolemia, or hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kane
                                                                                                                                                                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                            expression
                                                                                                                                                                                                                                                                                                                                                                                    oligonucleotide sequence is an antisense DNA oligo used to modulate the expression of the human LRH1 protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 3388; 909pp; English.
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                                                                       1607
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19
                                                                                                                                              l Similarity
17; Conserv
                                               AGCATGTCTTCTACTTCAG 1625
AGAATGTCTTCTAATTCAG 1
                                                                                                                                                                                                                                                                                                                                                                                of the
                                                                                                                                                                                                                                                                                                     BP; 7 A; 3 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cytidine
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cytidine nucleobases
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                                                                                                                                                                                                                                                                                                                                                                                human LRH1 protein
                                                                                                                                                                                      89.5%;
                                                                                                                                                                                                                              0.3%;
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                                                                                                                                                  0
                                                                                                                                                                                      Score 15.8;
Pred. No. 7
                                                                                                                                                  Mismatches
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ses are 5-methylcytidine."
                                                                                                                                                                                      .7e+02
                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                              Length 20
                                                                                                                                                      Indels
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                                                                                                                                                  0
                                                                                                                                                  Gaps
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RESULT 589
ADK80791
ID ADK807
XX ADK807
XX ADK807
XX CO-MAY
XX CO-
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache; seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2 MOS wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
         WO2004016754-A2
                                                                                                                                                     Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache;
                                                                                                                                                                                                                                                   Chimeric phosphorothioate oligonucleotide to target Nav1.3 #8125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 6 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 4; SEQ ID NO 1525; 417pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-FEB-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric phosphorothioate oligonucleotide to target Nav1.3 #1525
                                                                                                                                                                                                                                                                                                                       20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                ADK80791 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     infantile
                                                                                                                           intantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                     ADK80791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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Pred.
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No. 7
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7.7e+02;
2;
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RESULT 590
ADK75025/c
ID ADK750
XX A
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, diabetic neuropathy, trigeminal neuropathy, arthritic pain, acute pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache, seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothicate oligonucleotide with 2'MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target different regions of the human Navl.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Nav1.3; Analgesic; Nootropic; Neuroprotective; post-herpetic neuralgia; diabetic neuropathy; arthritic pain; migraine headache; infantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
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                                                              (PHAA ) PHARMACIA CORP
                                                                                                                            14-AUG-2002; 2002US-0403416P
                                                                                                                                                                                    14-AUG-2003; 2003WO-US025465.
                                                                                                                                                                                                                                                                                                                    WO2004016754-A2
                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric phosphorothioate oligonucleotide to target Nav1.3 #2359.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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Pred. No. 7.7e+02;
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RESULT 591
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Best Local S
Matches 17
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                                            New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associwith Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
                                                                                                                                                                                                                                                                                                                                                                                           Nav1.3; Analgesic; Nootropic; Neuroprotective; diabetic neuropathy; arthritic pain; migraine linfantile epilepsy; ataxia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric phosphorothioate oligonucleotide to target Navl.3 #7850.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compound targeted to a nucleic acid molecule encoding Nav1.3, useful for useful for treating a disease or condition associated with Nav1.3, e.g. pain, seizure disorder such as childhood seizure disorder, or ataxia.
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                                                                                                                                WPI, 2004-203785/19.
                                                                                                                                                                                                                               14-AUG-2002; 2002US-0403416P
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                                                                                                                                                                                              (PHAA)
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d molecule encoding Navl.3, where the antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A; 6 C; 4 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA; 20
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Pred. No. 7.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                             disease or condition associated such as childhood seizure
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                              ; post-herpetic neuralgia;
headache;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
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Claim

SEQ

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NO 7850;

417pp;

This invention describes a novel antisense oligonucleotides which

Example 15;

SEQ ID NO

51; 60pp;

English.

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RESULT 592
ADL34605/c
ID ADL346
XX ADL346
XX ADL346
XX ISIS a
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XX WPI; 2
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to an antisense compound targeted to a nucleic acid molecule encoding Navl.3, where the antisense compound specifically hybridizes with and inhibits the expression of Navl.3. The compound and composition are useful for treating a disease or condition associated with Navl.3, e.g. pain including but not limited to neuropathic pain, post-herpetic neuralgia, chronic pain, lower back pain, pain from burns, migraine headache, cluster headache, mild-to-moderate headache, seizure disorder such as childhood seizure disorder, including but not limited to neonatal or infantile epilepsy; or ataxia. The present sequence represents a chimeric phosphorothioate oligonucleotide with 2 MOE wings and a deoxy gap. Used during the antisense inhibition of human Navl.3 expression, the oligonucleotides are designed to target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antisense; inhibitor; hybridisation; human; phosphoinositide-3-kinase; regulatory subunit 4; p150; internucleoside linkage; phosphorothicate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine; infection; inflammation; tumour formation; hyperproliferative disorder cancer; Chediak-Higashi syndrome; metabolic disorder; immunomodulator;
                                                                      New antisense compound targeted to a nucleic acid molecule encoding phosphoinositide-3-kinase, regulatory subunit 4, p150, useful for treating cancer, Chediak-Higashi syndrome or a metabolic disorder.
                                                                                                                                                     WPI; 2004-282523/26.
                                                                                                                                                                                           Freier SM, Dobie KW
                                                                                                                                                                                                                                                                                      31-MAY-2002; 2002US-00160786
                                                                                                                                                                                                                                                                                                                         18-SEP-2003; 2003US-00667022.
                                                                                                                                                                                                                                                                                                                                                                                                    US2004063657-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADL34605;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                different regions of the human Nav1.3 RNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ISIS antisense oligonucleotide ISIS 206997.
                                                                                                                                                                                                                              (FREI/) FREIER S M
(DOBI/) DOBIE K W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1349 AAAAATTCACACAGCTGCT 1367
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No. 7.7e+02;
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This invention describes a novel antisense oligonucleotides which specifically hybridises to and inhibits the expression of human phosphoinositide-3-kinase, regulatory subunit 4, p150. The oligonucleotides comprises at least one modified internucleoside linkage, preferably a phosphorothioate linkage. It also comprises at least one modified sugar moiety, preferably a 2'-O-methoxyethyl sugar moiety. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense; inhibitor; hybridisation; human; phosphoinositide-3-kinase; regulatory subunit 4; p150; internucleoside linkage; phosphorothioate linkage; 2'-O-methoxyethyl sugar; 5-methylcytosine; infection; inflammation; tumour formation; hyperproliferative disorder; cancer; Chediak-Higashi syndrome; metabolic disorder; immunomodulator;
                                                                                                                                                                                                                                                                                     New antisense compound targeted to a nucleic acid molecule encoding phosphoinositide-3-kinase, regulatory subunit 4, p150, useful for treating cancer, Chediak-Higashi syndrome or a metabolic disorder.
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Pred. No. 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human; 88; interleukin-4 receptor; IL-4; interleukin-5 receptor; IL-5; CCR1; CCR3; Eotaxin-1; RANTES; MCP4; CD23; ICAM; VCAM; tryptase a; tryptase b; PDE4 A; PDE4 B; PDE4 C; PDE4 D; respiratory disease; lung disease; hyper-responsiveness; adenosine; adenosine A receptor; asthma; lung allergy; inflammation; inflammatory disease; airway inflammation; allergy; impeded respiration; cystic fibrosis; CF; chronic obstructive pulmonary disease; CP; allergic rhinitis; acute respiratory distress syndrome; pulmonary hypertension; lung inflammation; bronchitis; airway obstruction; bronchoconstriction.
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Pred. No. 7.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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Novel single or multiple target oligonucleotide anti-sense to e-initiation codon, intron of respiratory disease-relevant gene e-RANTES, MCP4, useful for prophylaxis or treating respiratory dis

disease

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Claim 2;

SEQ

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1707; 174pp;

English

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RESULT 595
AD031986/c
ID AD0319
XX AD0319
XX AD0319
XX AD0319
XX Cyclin
XX Cyclin
XX Antise
KW ANTise
RW Viral
XX VIRAL
XX INS
PN US2004
XX 31-JUL
XX 31-JUL
XX 31-JUL
XX 31-JUL
XX INS
PR 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ຊິຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄຄ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to oligonucleotides anti-sense to an initiation CC codon, coding region, 5' or 3' intron-exon junction, intron or region CC with 2-10 nucleotides of the 5'-end or 3'-end of a nucleic acid target CC chosen from a gene encoding interleukin (II)-4 receptor, interleukin (II) CC -5 receptor, CCR1, CCR3, Ectaxin-1, RANTES, MCP4, CD23, ICAM, VCAM, CC tryptase a tryptase b, PD84 A, PD84 B, PD84 C or PD84 D. The invention CC also relates to a method of screening a candidate compound that binds to come or more nucleic acid target(s) or expressed product(s), for the CC prevention and/or treatment of a respiratory or lung disease. The CC oligonucleotides are useful for reducing or inhibiting expression of a compound expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cyclin-dependent kinase 6, antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADO31986 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         allergic rhinitis, acute respiratory distress syndrome, pulmonary hypertension, lung inflammation, bronchitis, airway obstruction or bronchoconstriction. This sequence represents an oligonucleotide o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-JUL-2002; 2002US-00210802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-JUL-2002; 2002US-00210802
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2004087523-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       viral infection; apoptosis; ss; probe; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antisense therapy; cyclin-dependent kinase 6;
hyperproliferative disorder; cancer; bacteria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 7
                                                                                                                                                                                                                                                                         WPI; 2004-356241/33.
                                                                                                                                                                                                                                                                                                                                                    Freier SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3215 GACTGCAGCTGGTCAGCTG 3233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GACTGCAGCACGTCAGCTG 19
                                                                                                                                                                                                                                                                                                                                                                                                                               ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                    Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; 6 G; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.8;
Pred. No. 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cancer; bacterial infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .7e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 #83.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of the
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The invention relates to antisense oligonucleotides targeted to cyclindependent kinase 6, and which inhibit the expression of cyclin-dependent kinase 6. The antisense oligonucleotides are useful for treating a disease or condition associated with cyclin-dependent kinase 6, such as a hyperproliferative disorder (e.g. cancer), or conditions arising from bacterial or viral infections, or involving aberrant apoptosis. They are also useful in research and diagnostics for modulating the expression of cyclin-dependent kinase 6. The present sequence represents a cyclindependent kinase 6 antisense oligonucleotide. Note: Seqid 15-134 are also used in Tables 1 and 2 (page 30-34) but these sequences do not match seqid 15-134 of the seq list.
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                                                           ¤nnnnnnn x
                              Matches
                                    Query Match
Best Local (
                                                           Sequence 20 BP; 6 A; 1 C; 9 G; 4 T; 0 U; 0 Other;
              2318 CCATCATCTCCACCTTCTT 2336
19
                                    Similarity
CCACGATCTCCACCTTCTT
                             Conservative
                                    0.3%;
                             0
                                    Score 15.8; DB 1;
Pred. No. 7.7e+02;
                              Mismatches
                                           DB 1; Length 20
                              0;
                              Gaps
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0

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RESULT 596
AD031911
ID AD0319
XX AD0319
XX AD0319
XX 29-JUL
XX 29-JUL
XX Cyclin
XX antise
KW hyperp
KW viral
XX US2004
XX US2004
XX O6-MAY
XX 1-JUL
XX 31-JUL
XX ATSIE
PF 31-JUL
XX WPI; 2
XX WPI; 2
XX WPI; 2
XX WPI; 2
XX The ir
CC disease
CC disease
CC disease
CC disease
CC cyclir
CC cyclir
CC cyclir
CC cyclir
CC cyclir
CC cyclir
CC depend
CC cyclir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense therapy; cyclin-dependent kinase 6; hyperproliferative disorder; cancer; bacterial viral infection; apoptosis; ss; probe; human.
                                                                                                                                                                                                                                                                                                                             Freier SM, Dobie KW;
                                                                                                                                                                                                                                                                                                                                                                                    31-JUL-2002; 2002US-00210802.
                                                                                                                                                                                                                                                                                                                                                                                                                 31-JUL-2002; 2002US-00210802.
                                                                                                                                                                                                                                                                                                                                                                                                                                                06-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US2004087523-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cyclin-dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           29-JUL-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AD031911;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADO31911 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                         (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              kinase 6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              #8
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New compounds, particularly antisense oligonucleotides targeted to a nucleic acid encoding cyclin-dependent kinase 6, useful for treating cancer, bacterial/viral infection or conditions involving aberrant

WPI; 2004-356241/33.

Disclosure; SEQ ID NO 22; 68pp; English.

The invention relates to antisense oligonucleotides targeted to cyclindependent kinase 6, and which inhibit the expression of cyclin-dependent kinase 6. The antisense oligonucleotides are useful for treating a disease or condition associated with cyclin-dependent kinase 6, such as a hyperproliferative disorder (e.g. cancer), or conditions arising from bacterial or viral infections, or involving aberrant apoptosis. They are also useful in research and disposal to for modulating the expression of cyclin-dependent kinase 6. The present sequence represents a cyclindependent kinase 6 antisense oligonucleotide. Note: Seqid 15-134 are also used in Tables 1 and 2 (page 30-34) but these sequences do not match (page 30-34) but these sequences do |

New compounds, particularly antisense oligonucleotides targeted to nucleic acid encoding cyclin-dependent kinase 6, useful for treatic cancer, bacterial/viral infection or conditions involving aberrant

Disclosure; SEQ ID NO

97; 68pp; English

apoptosis.

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RESULT 597
ADN94856
ID ADN948
XX ADN948
XX ADN948
XX Human
XX Human
XX Homo
XX
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Query Match
Best Local Similarity
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                                                                                                                The invention relates to antisense oligonucleotides which are targeted to, and inhibit the expression of, a nucleic acid molecule encoding nidogen. The antisense oligonucleotides are useful for treating a disease or condition associated with nidogen, such as Chediak-Higashi syndrome. They are also useful in research and diagnostics for modulating the expression of nidogen. The present sequence represents a human nidogen antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                           New compounds, particularly oligonucleotides targeted to a nucleic acid encoding nidogen, useful for treating diseases associated with nidogen, e.g. Chediak-Higashi syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                        Sequence 20 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2004-389192/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-NOV-2002; 2002US-00300611
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human; Antisense Therapy; Gene Therapy; nidogen; Chediak-Higashi syndrome; ss; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human nidogen antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADN94856;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADN94856 standard;
                                                                                                                                                                                                                                                                                                              Example 15; SEQ ID NO 61; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2318 CCATCATCTCCACCTTCTT 2336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 4 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                        N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod_base= Other
/note= "Phosphorothioate
methylcytidines"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ₹,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16. .20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod_base= Other
/note= "2'-methoxyethyl
                                                                     Ą
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
0.3%;
89.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             _base= Other
                                                                     C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "2'-methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ი
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               G; 6 T; 0 U;
  Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.8;
Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
15.8; DB 1;
No. 7.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleotides'
                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleotides"
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RESULT 598
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ADN949
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ADN949
AC ADN949
A
  RESULT 599
ADP20520/c
ID ADP205
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AC AP-2 a
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Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to, and inhibit the expression of, a nucleic acid molecule encoding nidogen. The antisense oligonucleotides are useful for treating a disease or condition associated with nidogen, such as Chediak-Higashi syndrome. They are also useful in research and diagnostics for modulating the expression of nidogen. The present sequence represents a human nidogen antisense oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New compounds, particularly oligonucleotides targeted to a nucleic acid encoding nidogen, useful for treating diseases associated with nidogen, e.g. Chediak-Higashi syndrome.
cytostatic; AP-2-Inhik
AP-2 alpha associated
                                                                                Transcription factor AP-2 antisense oligonculeotide segid 67.
                                                                                                                                                                                                                                             ADP20520 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human; Antisense Therapy; Gene Therapy; nidogen; Chediak-Higashi syndrome; ss; probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12-AUG-2004
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                                                                                                                                    26-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to antisense oligonucleotides which are targeted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; SEQ ID NO 125; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2004-389192/36.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-NOV-2002; 2002US-00300611.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human nidogen antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADN94920
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                                                                                                                                                                                                                                                                                                                                                                               TGGTGCCAGCCCCATCCTG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Dobie KW;
  AP-2-Inhibitor-Alpha; AP-2 alpha; AP-2 associated disorder; hyperproliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 5 A; 5 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                    (first entry
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                                                                                                                                                                                                                                                DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.8; DB 1
Pred. No. 7.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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     alpha modulator;
disorder; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20;
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RESULT 600
AD085645/c
ID AD0856
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 specifically hybridises with a nucleic acid molecule encoding AP-2 alpha (19868 bp, SEQ ID NO: 4), and inhibits the expression of AP-2 alpha in cells or tissues comprising contacting the cells or tissues with (1); screening for a modulator of AP-2 alpha by contacting a preferred target segment of a nucleic acid molecule encoding AP-2 alpha with one or more candidate modulators of AP-2 alpha, and identifying one or more candidate alpha expression, which modulate the expression of AP-2 alpha; a disease state; and a kit or assay device comprising (1). The compound is useful for treating an animal having a disease or condition associated with AP-2 alpha, particularly a hyperproliferative disorder. The compounds may be used for diagnostics, therapeutics prophylaxis and as research reagents, or as tools in differential and/or combinatorial analyses to elucidate expression patterns of a portion or the entire complement of genes expressed within cells and tissues. This sequence represents a human transcription factor and a prise expressed and tissues of ligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New compounds targeted to nucleic acid molecules encoding AP-2 alpha and inhibits the expression of AP-2 alpha, useful for treating AP-2 alpha-associated disease or condition, particularly a hyperproliferative
                                                                                                                  CNS; zinedin modulator; zinedin; zinec neuronal disorder; zinedin modulator;
                                                                                                                                                                                                                                                                                                                                                                                            ADO85645 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention describes a compound (I) 8-80 nucleobases in length targeted to a nucleic acid molecule encoding AP-2 alpha. The compound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 15; SEQ ID NO 67; 58pp; English.
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                                                                                         antisense technology; ss.
                                                                                                                                                                                                            Human zinedin antisense oligonucleotide seqid
                                                                                                                                                                                                                                                                      26-AUG-2004
                                                                                                                                                                                                                                                                                                                                 ADO85645,
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                              Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ISIS-) ISIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transcription factor; antisense oligonucleotide; antisense technology;
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP;
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Pred. No. 7
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                                                                                                               zinedin associated disorder;
ator: human; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
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RESULT 601
AAQ75681/c
ID AAQ756
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AC AAQ756
XX
AC AAQ756
XX
OT 04-AUG

AAQ75681 standard; DNA; 21

ВP

04-AUG-1995 AAQ75681;

(first entry)

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CC to, and which specifically hybridizes with, a nucleic acid molecule care: inhibiting the expression of zinedin. Also described care: inhibiting the expression of zinedin in cells or tissues by contacting the cells or tissues with the compound so that expression of zinedin is inhibited; screening a modulator of zinedin by contacting a contacting the endidates of zinedin and identifying zinedin with one or more candidate modulators of zinedin, and identifying one or more condulators of zinedin expression which modulate the expression of zinedin a contacting a sample using any of the primers or probes given in the specification; a kit or assay device comprising the compound; and treating an animal a therapeutic or prophylactic amount of the compound so that expression zinedin is inhibited. The compount of the compound so that care useful for treating a disease or condition associated with zinedin, can useful for treating a disease or condition associated with zinedin, care useful for treating a disease or condition associated with zinedin, can be compound so that care useful for treating a disease or condition associated with zinedin, can be compound so that care useful for treating a disease or condition associated with zinedin, can be composited to the zinedin, and zinedin antisense or condition associated with zinedin, can be composited to zinedin, and zinedin antisense or condition. This sequence
                                                                                                                   Matches
                                                                                                                                                Query Match
Best Local :
                                                                                                                                                                                                                                       Sequence 20 BP; 3 A; 5 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 15; SEQ ID NO 81; 65pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.g. neuronal disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New compounds, particularly oligonucleotides targeted to a nucleic acid encoding zinedin, useful for treating diseases associated with zinedin,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10-DEC-2002; 2002US-00317270
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                                                                                                                   l Similarity
17; Conserv
CCTGCCTAGCCGTGGACCC 1
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/note= "OTHER= Phosphorothioate backbone. All cytidines
are 5-methylcytidines"
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mod_base= OTHER

mod_base= OTHER= 2'-O-Methoxyethyl
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                                                                                                                                                0.3%;
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                                                                                                                                                   Score 15.8;
Pred. No. 7
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RESULT 602
AAQ75730/c
ID AAQ757
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Best Local Similarity
Matches 17; Conserv
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                     WPI; 1995-018287/03.
                                                                                                                                                                                                                                04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 1 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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by digestion with restriction enzymes.
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                                                                                                                                                    Synthetic.
                                                                                                                                                                                    Analysis;
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                                                                16-APR-1993;
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                                          (NITE ) NIPPON
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                                                                                                                                                                                    gene expression; reverse transcription;
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cDNA and
                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                          TELEGRAPH & TELEPHONE
                                                                                    93JP-00112515
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gene
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expression -
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Pred.
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No. 7
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amplification
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                                                                                                                                                                                    primer; cDNA;
of.
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 mRNA
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Query Match Best Local S Matches 17

15; Conserv

0.3%; ilarity 89.5%; Conservative

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Mismatches

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Gaps

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Score 15.8; DB 1; Pred. No. 7.8e+02;

Length

21;

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RESULT 603
AAQ75728/c
ID AAQ75728;
XX AAQ75728;
XX AAQ75728;
XX AAAJ9sis;
XX Analysis;
YX 16-APR-19;
XX WPI; 1995
XX Disclosur
XX PT Analysis;
YT An
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Best Local Similarity
Matches 17; Conserv
                                                    A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-075798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. The method can be used to analyse gene expression rapidly and easily
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                                                                                                                                                                                                                                                                                                                                                                                                 Analysis of cDNA and gene expression -
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-018287/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   16-APR-1993;
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  BP; 3 A; 0 C; 0 G; 18 T; 0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           expression; reverse transcription; primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      93JP-00112515.
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2;
        0 Other;
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RESULT 604
AAQ75727;
ID AAQ75727;
XX
AC AAQ75727;
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AAQ75727;
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AAQ75727;
DT 04-AUG-1995 (first entry)
XX
C AAQ75727;
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AAA1ysis; gene expression, revents
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PN JP06303997-A.
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AAA1ysis; 93JP-00112515.
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PP 16-APR-1993; 93JP-00112515.
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DR WPI; 1995-018287/03.
XX
AAA1ysis of cDNA and gene expresion
CC Aamethod for the analysis of cD
CC A method for the analysis of cD
CC Cand using the aggregate of mRNA
CC transcription primer; (b) diges
CC the double-stranded cDNAs with
CC electrophoresing the digested a
CC method can be used to analyse g
XX
Sequence 21 BP; 2 A; 0 C; 1 G;
RESULT 605
ADG77640
ID ADG776
XX ADG776
XX ADG776
XX II-MAR
XX Canine
XX Canine
XX Geneti
XW Geneti
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Best Local S
Matches 17
                                                                                                         genetic disease; genetic trait; dog; carrier of recessive disease; copper toxicosis; CT; canine genome map; breed-specific profile; DNA fingerprint; dog identification; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A method for the analysis of cDNA comprises (a) preparing an aggregate (double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The electrophoresing the digested aggregate of cDNAs in separate lanes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Analysis of cDNA and gene expression - by amplification of mRNA followed by digestion with restriction enzymes.
                                                                    Canis familiaris.
                                                                                                                                                                                                        Canine disease marker-related PCR primer 484.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 2 A; 0 C; 1 G; 18 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 8; 11pp; Japanese.
                                                                                                                                                                                                                                                    11-MAR-2004
                                                                                                                                                                                                                                                                                                 ADG77640;
                                                                                                                                                                                                                                                                                                                                         ADG77640 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Reverse transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5390 ATTAAAAAATACAAAAAA 5408
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTAAAAAAAAAAAAAAA 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene expression; reverse transcription; primer; cDNA; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     be used to analyse gene expression rapidly and easily
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                                                                                                                                                                                                                                                    (first entry)
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Pred. No. 7.
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RESULT 606
AAX32871/c
ID AAX328
XX AAX328
XX AAX328
XX CAAX328
XX Triple
XX Triple
XX Inhibi
XX Synthe
OS Hepati
XX WO9920
XX Synthe
PD 29-APR
XX WO9920
XX 29-APR
XX 21-OCT
XX 21-OCT
XX 21-OCT
XX X1 Lu C;
PA (SHAN-
XX Lu C;
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                      27-AUG-2003
20-MAR-2003
28-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to novel oligonucleotide PCR primers which may be used to identify markers associated with genetic diseases and traits in dogs, in particular to disgnose genetic diseases that are not phenotypically visible and to identify carriers of recessive diseases. A specific application is diagnosis of copper toxicosis (CT). The invention can also be used to create a genetic map of the canine genome; to generate breed-specific profiles; to establish paternity and to identify dogs from DNA fingerprints. The method provides rapid analysis of the target sequences from only a small sample of DNA. Diagnosis can be done at any time in the dog's life. The present sequence is that of a PCR primer of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New oligonucleotide primers for diagnosis of genetic diseases and train dogs - amplify specific regions of the genome containing microsatellite repeats, especially for diagnosing copper toxicosis and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UNMI)
                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                               Triplex-forming oligonucleotide; TFO; promoter region; pre-S gene; inhibition; hepatitis B virus; HBV adr subtype; DR region; ss.
                                                                                                                                                                                                                                                                                                                                                                   AAX32871;
                                                                                                                                                                                                                                                                                                                                                                                              AAX32871 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 21 BP; 6 A; 6 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 15; 40pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           carriers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Brewer GJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-FEB-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-AUG-1997.
                                                                                                  19-OCT-1998;
                                                                                                                               29-APR-1999
                                                                                                                                                          WO9920641-A1
                                                                                                                                                                                       Hepatitis B virus.
                                                                                                                                                                                                                                                                         TFO B13 sequence.
                                          (SHAN-) SHANGHAI INST BIOCHEMISTRY CHINESE ACAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2406
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17; Conserv
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UNIV MICHIGAN STATE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                      (revised)
(revised)
(first en
                                                                       97CN-00106667.
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                                                                                                    98WO-CN000248
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                                                                                                                         entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.8; DB 1
Pred. No. 7.8e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
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RESULT 607
AAX57097
ID AAX570
XX AAX570
XX AAX570
XX CAQ2;
XW CCQ2;
XW CCQ2
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    and TFO B11, B12 and B15 (AAX32868-870) can bind with DR region of HBV. The oligonucleotides are useful for inhibition of HBV and as drug in treatment of hepatitis B. Since the length of the oligonucleotides can be suitably increased, the stability and specificity of the formed triplex DNA with 2 similar homopoly purine/homopoly pyrimidine fragments are higher. Triplex formation is specifically targeting on the HBV gene expression, DNA replication and reproduction, or to produce (DNA)2:RNA hybrid triplex with target sequence of RNA in stopping RNA reverse transcription, so there is little effect on the human cells. Such oligionucleotides are chemically modified by 3'-terminal monophosphorylation, leading to more significant inhibition due to their higher stability, and the degradation products of the modified oligonucleotides are not toxic to the body. (Updated on 20-MAR-2003 to correct DR field.) (Updated on 27-AUG-2003 to correct OS field.)
                                                                                Claim 63; Page 145; 195pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-MAY-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rolandic epilepsy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          KCNQ2; KCNQ3; human; murine; potassium channel; diagnosis;
benign familial neonatal epilepsy; BFNE; juvenile myotonic
rolandic epilepsy; mutant; treatment; screening; epilepsy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention provides triplex-forming oligonuclectides (TFO) modified derivatives. TFO B1-B5 (AAX32862-866) can bind with t region of pre-S gene in inhibition of hepatitis B virus (HBV) and TFO B11, B12 and B15 (AAX32868-870) can bind with DR region of the provided by the control of the co
                                                                                                                                                                 Nucleic
                                                                                                                                                                                                                                              WPI; 1999-312938/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-OCT-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9921875-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX57097 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 12; 39pp; Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hepatitis B virus
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                                                                                                                                                                                                                                                                                                                       NA,
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                                                                                                                                                                 acid
                                                                                                                                                                                                                                                                                                                       Leppert MF,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5 A; 0 C; 16 G; 0 T; 0 U; 0 Other;
                                                                                                                                                         encoding potassium channels KCNQ2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             screening;
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                                                                                                                                                                                                                                                                                                                       Charlier
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Pred. No. 7
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                                                                                                                                                             ω
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; prognosis;
c epilepsy; JME;
; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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lr subtype
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invention describes novel human and mouse potassium channel proteins

Antisense oligonucleotides (ASO) for inhibiting a tumour necrosis factoralpha (TNF-alpha) gene in an animal, preferably a human, comprise 12-50 nucleotides, 90% of which are complementary to a region of mRNA containing a GGGA sequence motif. The ASO is used to inhibit expression of a gene in an animal and for treating the animal when afflicted with a disease or disorder characterised by the presence of an mRNA from a gene containing a GGGA motif. The ASO are specifically targeted to a GGGA sequence motif found in mRNA from a gene. A study of known ASO has shown that at least half of the most efficacious ASO's contain one or more TCCC

Generation of antisense oligonucleotides - GGGA motif found in mRNA sequences.

by specifically targeting a

Example 1; Page 32; 55pp; English.

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RESULT 608
AAXO9047
ID AAXO90
XX AAXO90
XX Tumoux
XX Tumoux
XX Tumoux
XX Inhibi
XX Synthe
OS Rattus
XX WO9901
XX ATTUS
PM 03-JUJ
XX WPI;
XX WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CC KCNQ2 and KCNQ3. Detecting mutations in sequences that encode KCNQ2 or CC KCNQ3, or the loss of one copy of these genes, is used for diagnosis and CC prognosis of benign familial neonatal epilepsy (BFNE), juvenile myotonic CC epilepsy (JME) or rolandic epilepsy (RE). Cells (or transgenic animals) CC that express wild-type or mutant KCNQ2 or 3 (also the proteins themselves CC in cell-free form) are used to screen for agents that can be used to creat or prevent these forms of epilepsy. Fragments of the encoding CC mutations or for isolation of related sequences, while the complete CC sequences may be used in gene therapy to provide wild-type proteins. CC Antibodies specific for mutant or wild-type proteins are used as CC diagnostic reagents and for drug screening. The KCNQ2 and 3 proteins are useful in rational design of drugs and therapeutically (in replacement CC therapies). The forms of epilepsy associated with mutations in KCNQ2 and CC sequences can now be diagnosed early (before symptoms are manifest), CC and better treatment options will be available. AAX57074-X57139 are primers used in the method of the invention
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tumour necrosis factor alpha; TNF-alpha; antisense oligonucleotide; ASO: inhibition; expresssion; treatment; disease; disorder; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                    WPI; 1999-105767/09.
                                                                                                                                                                                                                                                                                                                                                                             03-JUL-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9901139-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX09047 standard; DNA; 21 BP
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                                                                                                                                                                                                                                                                                                                                                                                                            02-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JAN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rattus rattus.
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                                                                                                                                                                                                                                                                                                       Israel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            necrosis factor alpha antisense oligonucleotide
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Pred. No. 7.
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Best Local
                         The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the diagnosis of atherosclerosis, myoccardial infarction, coronary heart disease, stroke, peripheral vascular disease, venous thrombosembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPS) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                              Nucleic acids comprising single nucleotide polymorphisms, useful in applications such as forensics, paternity testing, medicine, genetic analysis and phenotype correlations to diseases such as diabetes and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; variant thrombospondin 1; variant thrombospondin 4; SNP; polymorphism; vascular disease; coronary artery disease; forensics; myocardial infarction; atherosclarosis; stroke; venous thromboembolism; pulmonary embolism; paternity test; ds.
                                                                                                                                                                                                                                                                                                                                      Lander ES,
                                                                                                                                                                                                                                                                                                                                                                  (MILL-)
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26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human gene single nucleotide polymorphism #1207.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        motifs. This ASO was designated TJU-2698 and corresponds to the 3' untranslated region of the primary transcript of rat
                                                                                                                                                                                                      Example; Page 133; 242pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-MAR-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Variation
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                                                                                                                                                                                                                                                                                                                                     Gargill M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /standard_name= "single nucleotide polymorphism"
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Pred. No. 7.8e+02;
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TNF-alpha
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Sequence 21 BP; 4 A;

10 C; 4 G; 3 T; 0 U; 0 Other;

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                                                     Query Match
Best Local S
Matches 17
                                                                                                                                 The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothioate backbone
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                Vaccinating against tumors, infectious diseases, allergies and using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-SEP-1999; 99US-0156113P
27-SEP-1999; 99US-0156135P
23-AUG-2000; 2000US-0227436P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothicate; 85.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAF99580 standard; DNA; 21
                                                                                                             Sequence 21
                                                                                                                                                                                                                                                                                                                                                      Claim 101; Page 53; 338pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-SEP-2000; 2000WO-US026383.
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                                                     l Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Schetter C,
                                                                                                              BP; 0
                                                      Conservative
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99US-0156135P.
                                                                                                             A; 7 C; 7 G; 7 T; 0 U;
                                                                  0.3%;
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Pred. No. 7.8e
0; Mismatches
 19
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                                                                   Score 15.8; DB 1
Pred. No. 7.8e+02
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                                                        Mismatches
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RESULT 611
AAF76158
ID AAF761
      The invention relates to a transgenic immunodeficient recipient mouse CC which is capable of supporting the growth of donor cells. In the mouse, CC both alleles of a gene activated in early lymphocyte development are CC disrupted, causing it to lack mature B and T cells. In particular, both CC alleles of the recombination activation gene-2 (RAG-2) gene are CC disrupted, which in turn prevents VDJ recombination. The mouse also CC comprises donor (e.g., human) specific transgenes encoding the cytokines CC interleukin-7 (IL-7), stem cell factor (SCF), leukaemia inhibitory factor (LIF), granulocyte macrophage-colony stimulating factor (M-CSF), and IL-6, which enable it CC macrophage-colony stimulating factor cells. In another embodiment CC transgene has naturally linked DRAB and Gonor cells. In another embodiment CC transgene has naturally linked DRAB in DRAB molecule, where the transgenic mouse may be used as a model for determining the allergenicity of non-donor, e.g., non-human, macromolecules; to determine the effect compounds have on a human immune system; to generate fully human polyclonal or constitute of the monoclonal antibodies will raise a response in a human immune system; to entermine whether
pathogens and other immunomodulatory compounds; and to determine the factors involved in regulating the development and function of human haematopoietic cells. The transgenic mouse supports the functional properties of human haematopoietic cells, unlike previous animal models which produce functionally impaired haematopoietic cells or are immunologically dysfunctional. In addition the transgenic mouse provides a unique model system which supports T cell development in a manner which more closely resembles normal ontogeny, as they possess CD4+ T cells in the periphery that exhibit MHC-restricted antigen- specific responses. Sequences AAF76133-AAF76192 represent human cytokine PCR primers used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 granulocyte macrophage-colony stimulating factor; macrophage-colony stimulating factor; human MHC class II; DR3; major histocompatibility complex; allergenicity determination; human monoclonal antibody generation; haematopoietic cell development;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 08-MAR-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          specific transgenes encoding cytokines.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New transgenic mice, useful as non-human mammalian models of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-169001/17.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    lymphocyte deficient; human cytokine; interleukin; IL-7; IL-6
stem cell factor; leukaemia inhibitory factor; GM-CSF; M-CSF;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Transgenic mouse; immunodeficient; tissue recipient;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human oncostatin M (OM) PCR primer,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immune system animal model; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              comprise recombination activation gene mutations and donor
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The invention relates to Detecting a collagen gene alteration associated with a pathological condition in a human subject by obtaining from the consecutive nuclectides of the segment of the COLIA1 gene extending in the 5' to 3' direction from 78 nucleotides of intron 27 located adjacent exon 28 through the 3' end of intron 51, where the portion contains an collection content and a first and second site, determining the sequence of the portion with the corresponding consensus sequence of the COLIA1 gene where a difference between the sequence of the portion and the consensus sequence comparing the sequence of the portion with the extreme the sequence of the COLIA1 gene where a difference between the sequence of the COLIA1 gene where a difference conteven the sequence of the COLIA1 gene where a difference conteven the sequence of the COLIA1 gene where a difference conteven the sequence of the COLIA1 gene where a difference conteven the sequence of the COLIA1 gene where a difference conteven the sequence of the COLIA1 gene where a difference conteven the sequence of the COLIA1 gene such as options is useful for context and the consensus sequence indicates the presence of the collagen alteration in the subject. The method is associated with an altered collagen gene such as osteoporosis, multiple epiphyseal dysplasia, osteogenesis imperfecta, shortness of stature and low bone density. Identification of an abnormality in a collagen gene is also useful for designing a therapeutic nucleotide or gene therapy agent which can be administered to the subject to correct or alleviate the correct or alleviate the mich can be administered to the subject to correct or alleviate the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detecting collagen gene alteration, useful for diagnosing osteoporosis, multiple epiphyseal dysplasia, osteogenesis imperfecta, shortness of stature and low bone density in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; collagen; COL1A1; COL1A2; COL9A1; COL9A2; COL9A3; ss; osteoporosis; multiple epiphyseal dysplasia; osteogenesis imperfe shortness of stature; low bone density; gene therapy; PCR primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Westerhausen Larson A,
Ala-Kokko L, Annunen S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Prockop DJ,
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(UYJE-) UNIV JEFFERSON THOMAS.
(UYOU-) UNIV OULU.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-DEC-1991;
13-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-JUL-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fig 25; 617pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Spotila LD, Deltas CD, Sereda L;
Larson A, Pack M, Colige A, Early
Annunen S, Pihlajamaa T, Vuoristo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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94US-00212322.
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Pred. No. 7.
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Paassilta P;
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                                                                                                                                                                                              The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, reconstants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; rheumatroid arthritis; psoriasis; diabetic retinopathy; retinopathy rematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coding and non-coding sequences of any of the COL1 or COL9 genes. Therefore the method can be used to detect collagen gene alterations which affect either the primary sequence of a collagen protein chain, splicing of the mRNA encoding such chains or regulation of expression the genes encoding such chains. The present sequence is a PCR primer which amplifies a nucleic acid from a collagen gene of the invention
                                                                                                 rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-566690/60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2; Page 33; 276pp; English
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                                       invention
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                                                                       present sequence
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Pred. No. 7
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                                                                       is an antiangiogenic nucleic
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RESULT 614
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ID ABL388
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YT Modiff
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Matches
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                                                                                The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, oseophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, mon-Hodgkin's lymphoma, melanoma, mylong oral cavity cancer, ovarian cancer, stomach cancer, testicular cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression o cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 156; 312pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      developing cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-JUN-2000; 2000US-0213346P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-JUN-2001; 2001WO-US020154.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-APR-2002
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                                                             exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Weiner G,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2641 CTGCAGCTGCTGCAGC 2659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2002-154611/20
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CIGCIGCIGCIGCIGCIGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hartmann
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/mod_base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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Pred. No. 7.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        e expression of or at risk of
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Query Match Best Local Similarity Thes 17; Conserve

Conservative

0

Mismatches

Indels

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0.3%;

Score 15.8; DB 1 Pred. No. 7.8e+02

DB 1;

Length

21;

Sequence 21

BP; 0

A; 7 C; 7 G; 7 T; 0 U; 0 Other;

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RESULT 615
AAL44216/c
ID AAL442
XX AAL442
XX Human
XX Human
XX Human
XX Homo &
XX Homo
ABK10202
ID ABK1
XX
AC ABK1
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AC ABK1
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DT 21-W
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DS Doub
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                                                                                                                                                                                                                                                                                                                           RESULT 616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention comprises a method for detecting single nucleotide polymorphisms (SNPs) in genes associated with human bronchial asthma. The method specifically refers to detecting polymorphisms in the gene encoding human I-kappa beta associated protein (IXAP). The invention also comprises primers and probes for use in the method of the invention. The method of the invention is useful for analysis of the etiology of bronchial asthma (especially infant bronchial asthma). The present DNA sequence represents a human I-kappa beta associated protein gene PCR
                              Single stranded DNA isolation; DNA purification; CTG repeat; ds.
                                                                                                                                                                                                           ABK10202;
                                                                                                                                                                                                                                                                     ABK10202 standard; DNA; 21 BP
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                                                                                  Double stranded DNA isolation (CTG) 7 repeat sequence.
                                                                                                                                                 21-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detection of specific single nucleotide polymorphisms in human I-kappa
beta associated protein for analysis of etiology of bronchial asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.8;
Pred. No. 7
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RESULT 617
ACC42594
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AC ACC425
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AC ACC425
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DT 26-AUG
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DE Human
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                                                                     Query Match
Best Local S
Matches 17
                                                                                                                                                                          nucleic acid that is a polymerase chain reaction (PCR) amplification product from a pool of related or unrelated sequences in high yield for subsequent use. The method also permits capture and/or recovery of the first strand of a double-stranded target nucleic acid from biological samples or other samples containing large molecule contaminants. The present sequence represents a double stranded (CTG) 7 DNA molecule used to isolate double stranded DNA molecules in an example of a similar method to that of the invention
                                                                                                                                                                                                                                                                                                                    This invention relates to a novel method for isolating one strand of double-stranded target nucleic acid. The method comprises contacting a double stranded target DNA molecule with a competitor oligonucleotide capable of hybridising to the first strand of the double stranded molecule. The method is performed under conditions in which the first strand dissociates from the second and hybridises with the competitor oligonucleotide to form a heteroduplex. The method of the invention is useful for separating a strand from a double-stranded target nucleic acid. The method is rapid, efficient and specific for isolating a single strand from a double-stranded nucleic acid. Because the method is advantageously used to purify a first strand from a double-stranded advantageously used to purify a first strand from a double-stranded
                                                                                                                                          Sequence 21 BP; 0 A; 7 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 12; 61pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Isolating one strand of double-stranded nucleic acid, by contacting double stranded nucleic acid having first and second strands with competitor oligo to form first strand-oligo complex and isolating the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chiesa C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-AUG-2000; 2000US-0222686P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-JUL-2001; 2001WO-US022782.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           repeat_unit
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                      2641 CTGCAGCTGCTGCTGCAGC 2659
                                                                     l Similarity
17; Conserv
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                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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l. .3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note= "CTG type repeat"
                                                                                      89.5%;
                                                                                                        0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Egholm M;
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                                                                     0; Mismatches
                                                                                       Score 15.8; DB 1; Length 21; Pred. No. 7.8e+02;
                                                                       Indels
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                                                                       Gaps
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Human oncostatin M, OM, PCR primer OSM3R1

26-AUG-2003 ACC42594;

(first entry)

ACC42594 standard; DNA; 21

ВP

<u>.</u>

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RESULT 618
ACH03118
ID ACH031
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ACH031
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a new transgenic mouse, which comprises a disruption in both alleles of a gene such that lymphocyte maturation does not occur and exogenous cytchines. The cytchines are selected from: interleukin-3 (IL-3), interleukin-6 (II-6), interleukin-7 (IL-7), macrophage-colony stimulating factor (M-CSP), granulocyte-colony stimulating factor (GM-CSP), pranulocyte-colony stimulating factor (GM-CSP), leukaemia inhibitory factor (IIF) and oncostatin M (OM). The gene disruption is in a gene that modulated VDJ recombination e.g. a RAG gene. The gene is disrupted by insertion of a transgene comprising major histocompatibility complex (MHC) Class II DR3 and DQ2 genes. The transgenic mouse is useful as a model of human disease to assess efficacy of therapeutic or prophylactic transgenic mouse is also useful for supporting donor haematopoletic stem cells or facilitating production of donor-specific functional immunity.

PCR primers ACC42571-ACC42639 were used to generate the transgenic mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
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Matches 17
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                                                                                                                             Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic context dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New recipient mammal, preferably a mouse, useful as a model of human disease to assess efficacy of therapeutic or prophylactic treatments, for facilitating production of donor-specific functional immunity.
                                                                                                                                                                                                                                                                                                                                                                                                                        ACH03118
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      US2003050268-A1
                                                                 Synthetic
                                                                                                                                                                                                                                                                                           Immunostimulatory nucleic acid #753.
                                                                                                                                                                                                                                                                                                                                                         25-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GEMV)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GATGCTGAGAAGGGGAGAG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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Pred. No. 7.8e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treating non-allergic inflammatory diseases, such as psoriasis, allergic contact dermatitis, latex dermatitis or inflammatory be disease by administering an immunostimulatory nucleic acid.
                                                                                 WPI; 2003-657977/62
                                                                                                                                                                                   (BRAT/)
(PETE/)
(FOUR/)
                                                                                                                                                                                                                                                                                          03-FEB-2000; 2000US-0179991P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ds; allergy; asthma; poly-G nucleic acid; aerosol hypo-responsive subject; immunostimulatory.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunostimulatory nucleic acid #696.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sequence represents an immunostimulatory nucleic acid
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                                                                                                                                                                                   PETERSEN D M.
FOURON Y.
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Pred. No. 7.8e+02;
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Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.

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                   patient. The method comprises determining the level of oestrogen-related receptor alpha (ERRalpha) expression in breast cancer cells, where a high level indicates a poor prognosis and a low level indicates a more favourable prognosis. Also disclosed is a method of categorising breast cancer patients based on ERRalpha status by determining the expression level of ERRalpha in the breast cancer cells of the breast cancer patient, a method of identifying a human subject for further breast cancer examination by determining whether a mutation exists in the
                                                                                                                                                                                                                                                                                                                                                                          Determination of prognosis of breast cancer patient comprises determining level of estrogen-related receptor alpha expression in breast cancer cells where high level indicates poor prognosis and low level indicates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Breast cancer prognosis; human; oestrogen-related receptor ERRalpha; breast cancer cell; ERE-dependent transcription;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mertz JE,
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(JOHN/)
(KRAU/)
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26-FEB-1998; 98US-00031250
20-JAN-2000; 2000US-00488730
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADG45359;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                               The present invention relates
                                                                                                                                                                                                                                                                                                    Example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-AUG-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ARIA/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       invention relates to a method of treating or preventing allergy or hma which comprises administering to a subject a poly-G nucleic aci
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2641
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                                                                                                                                                                                                                                                                                                                                                      favorable
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MERTZ J E.
JOHNSTON S D.
KRAUS R J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTGCAGCTGCTGCTGCAGC 2659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21 BP; 0 A; 7 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CIGCIGCIGCIGCIGC 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Johnston SD,
                                                                                                                                                                                                                                                                                                    SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                    prognosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA;
     by determining whether a mutation ethod of treating breast cancer by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -00031250
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                                                                                                                                                                                                                                                                                               7; 76pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21
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  treating
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                                                                                                                                                                                                                                            to methods of prognosis of a breast cancer
                                                                                                                                                                                                                                                                                                 English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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No. 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ariazi EA;
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exists in the determining w
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR; p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                       where a high
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     primer;
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ERRalpha is an activator or repressor of ERE-dependent transcription in the breast cancer cells, and decreasing ERRalpha activity in the breast cancer cells if ERRalpha is an activator and increasing ERRalpha activity in the breast cancer cells if ERRalpha is a repressor, and a method of identifying an agent that modulates ERRalpha expression. The methods of the invention are useful determining prognosis of a breast cancer patient. The invention identifies more biomarkers and treatment targets for breast cancer. The biomarker identified can predict prognosis and help patients make treatment choices. The present sequence represents a PCR primer used in the examples of the present invention.
Sequence 21
BP; 5 A; 5 C; 4 G; 7 T; 0 U; 0 Other;
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Query Match
Best Local S
Matches 17
             2720
N
                                  Similarity
             AAGTGATGGCCCATTTCGA 2738
                           Conservative
                                  0.3%;
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•
                                  Score
Pred.
20
                            Mismatches
                                  No. 7.
                                   .8e+02
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                                           <u>;</u>
                                         Length
                             Indels
                                           21;
                             <u>.</u>
                             Gaps
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ADJ13145 standard; DNA; 21
Human
                               20-MAY-2004
                                                             ADJ13145
DNA
probe used to immobilise CpG methylated DNA SeqID 272.
                             (first entry)
                                                                                              BP.
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probe; sapiens. ss; chemical modification; methylation; array; suppressor; p16; human; H69; H1618. suppressor; p16; CpG island

27-JUN-2001; 2001US-0301370P 27-JUN-2002; 2002US-00184085 14-AUG-2003 US2003152950-A1 (GARN/) GARNER H R.

(LUEB/) (NINN/) HR,) MINNA J D.) LUEBKE K J.) BALOG R P. Minna JD, Luebke ፭ Balog RP.

WPI; 2003-874843/81.

Analysis of chemical modification be analyzed, treating DNA with chemical reagents that result in ferent base sequences, and determining entermining that result in

1; SEQ ID NO 272; 210pp; English

human a human This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the

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RESULT 622
ADJ13037/c
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                            Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                            This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (GARN/) GARNER H R.
(MINN/) MINNA J D.
(LUEB/) LUEBKE K J.
(BALO/) BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                          Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Garner HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                probe; 88; chemical modification; methylation; array; CpG island;
tumour suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human DNA probe used to immobilise CpG methylated DNA SeqID 164.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21 BP; 4 A; 12 C; 0
                                                                                        Sequence 21 BP; 3 A; 12 C; 0 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                               Example 1; SEQ ID NO 164; 210pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-JUN-2001; 2001US-0301370P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADJ13037;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADJ13037 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-874843/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-2002; 2002US-00184085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-AUG-2003.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2436 GGATGAGAAGGGGAGAGGT 2454
 2436 GGATGAGAAGGGGAGAGGT 2454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 GGATGAGAGTGGGAGAGGT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Minna JD,
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                             Conservative
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                                        0.3%;
89.5%;
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                             0; Mismatches
                                        Score 15.8; DB 1;
Pred. No. 7.8e+02;
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Pred. No. 7.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ξ,
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                                                          DB 1;
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                             2; Indels
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                             Gaps
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RESULT 623
ADJ13036/c
ID ADJ130
XX
AC ADJ130
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Best Local S
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(MINN/) MINNA J D.
(LUEB/) LUEBKE K J.
(BALO/) BALOG R P.
                                                                                                                                                                  This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene p16 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                     Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                         Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-874843/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Garner HR, Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-JUN-2002; 2002US-00184085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            probe; ss; chemical modification; methylation; array; CpG island;
tumour suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human DNA probe used to immobilise CpG methylated DNA SeqID 163.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-MAY-2004
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                                                                                                                                                        invention.
                                                                                                                                                                                                                                                                                                                                                                                             Example 1; SEQ ID NO 163; 210pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-JUN-2001; 2001US-0301370P
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                             2436
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21
                                                          1 Similarity
17; Conserv
 GGATGAGAGGAGGAGGT 3
                             GGATGAGAAGGGAAGGT 2454
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                                                                                                                       BP; 3
                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                       P
                                                                         0.3%;
                                                                                                                       12 C; 0 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Luebke
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                                                          0
                                                         Score 15.8; DB 1
Pred. No. 7.8e+02
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Balog
                                                                                        DB 1; Length 21;
                                                              Indels
                                                            0
                                                            Gaps
                                                            0
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ADJ13072/c ID ADJ130 XX AC ADJ130

ADJ13072 standard; DNA; 21

BP

ADJ13072;

RESULT 624

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RESULT 625
ADJ13144/c
ID ADJ131
XX
AC ADJ131
XX
PT 20-MAY
YX
PT 20-MAY
YX
WX
DE Human
XX
CR
KW probe;
KW probe;
KW tumoux
XX
XX
OS Homo 8
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                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                        modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                              probe; ss; chemical modification; methylation; array;
tumour suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      probe; ss; chemical modification; methylation; array; CpG island;
tumour suppressor; p16; human; H69; H1618.
                                                                               Human DNA probe used to immobilise CpG methylated DNA SeqID 271.
                                                                                                                                                                                  ADJ13144 standard;
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21 BP; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to a novel method for analysing chemically
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; SEQ ID NO 199; 210pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-874843/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Garner HR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (GARN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-JUN-2001; 2001US-0301370P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-JUN-2002; 2002US-00184085
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2003152950-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human DNA probe used to immobilise CpG methylated DNA SeqID 199.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-MAY-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                               invention.
                                                                                                                                                                                                                                                                                                2436 GGATGAGAAGGGGAGAGGT 2454
                                                                                                                                                                                                                                                                 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GARNER H R.
) MINNA J D.
) LUEBKE K J.
) BALOG R P.
                                                                                                                                                                                                                                                                 GGATGAGAGGCGGAGAGGT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Minna JD,
                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                               A; 12 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                              0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Luebke
                                                                                                                                                                                  21
                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                              Score 15.8; DB 1;
Pred. No. 7.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ኟ
                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RP;
                                                                                                                                                                                                                                                                                                                                2
                                                                                                                                                                                                                                                                                                                                                           Length 21;
                                                                                                                                                                                                                                                                                                                                  Indels
                                                 CpG island
                                                                                                                                                                                                                                                                                                                                0
                                                                                                                                                                                                                                                                                                                                  Gaps
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RESULT 626
ADJ13073/c
ID ADJ130
XX ADJ130
XX DJ130
XX PT 20-MAY
XX PT UMMAN
XX PTODE;
KW EUMOUZ
XX EUMOUZ
XX US2003
XX US2003
XX US2003
XX 11-AUC
XX PF 27-JUN
XX PR 27-JUN
XX 27-JUN
XX 27-JUN
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many pottantial sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Analysis of chemical modification of DNA involves obtaining sample of
to be analyzed, treating DNA with chemical reagents that result in
different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                         Human DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Garner HR, Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-AUG-2003
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                                                                                                                                                                                                                                                                                       ADJ13073
                                                                                                                                                                                                                                                                                                                    ADJ13073 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-874843/81.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GARN/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               27-JUN-2002; 2002US-00184085.
                 27-JUN-2001; 2001US-0301370P
                                                27-JUN-2002; 2002US-00184085
                                                                               14-AUG-2003
                                                                                                              US2003152950-A1
                                                                                                                                                                                                                                                        20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                              2436 GGATGAGAAGGGGAGAGGT 2454
                                                                                                                                                                                                                                                                                                                                                                                                 21
                                                                                                                                                                         ss; chemical modification; methylation; array; CpG island
suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LUEBKE K
BALOG R P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GARNER H R.
MINNA J D.
                                                                                                                                                                                                                         probe used to immobilise CpG methylated DNA SeqID 200.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 4 A;
                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15.8; DB 1;
Pred. No. 7.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           G; 5 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Balog
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 21;
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RESULT 627
ADJ31733
ID ADJ317
XX ADJ
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in different base sequences depending upon the presence or absence of the modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analysis of chemical modification of DNA involves obtaining sample of to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
New compounds, particularly antisense nucleic acid encoding an amyloid beta
                                                                                                                                                                                                                                                                                                                                                                                                                         US2003232435-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Amyloid beta protein precursor; neurodegenerative disorder;
Alzheimer's disease; apoptosis; diagnosis; therapy; human; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human amyloid beta precursor DNA amplifying reverse PCR primer
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(LUEB/)
(BALO/)
                                                                                          WPI; 2004-061283/06
                                                                                                                                                                                                                                                                                                                 14-JUN-2002; 2002US-00173208
                                                                                                                                                                                                                                                                                                                                                                        18-DEC-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADJ31733 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1, SEQ ID NO 200, 210pp; English.
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                                                                                                                                                                                                                                                         14-JUN-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2436 GGATGAGAAGGGGAGAGGT 2454
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BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GARNER H R.
                                                                                                                                                                                                     SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GGATGAGAGCGGGAGAGGT 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                         2002US-00173208
                                                                                                                                                                                                     PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; 12 C; 1 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21
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Pred. No. 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Balog RP;
  oligonucleotides targeted protein precursor, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .8e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 21;
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        for
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RESULT 628
ADL70482
ID ADL704
XX ADL704
XX ADL704
XX ADL704
XX ANA insuli
KW Insuli
KW Insuli
KW Insuli
KW INSUli
FT MODIFI
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XX WOOLOG
YX WOOLOG
YX UNDRED
PT GONOS
XX UNBER
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XX UNBR
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PT GISSE
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PT TEANS
PT HESE
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Matches
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                                                                                                                                                                                                                                                                                                Jansen B,
Gonos ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The presentinvention is directed to antisense oligonuclectides targetted to a nucleic acid encoding amyloid beta protein precursor and which modulates the expression of amyloid beta protein precursor. The invention is useful for treating a disease or condition associated with amyloid beta protein precursor such as a neurodegenerative disorder e.g. Alzheimer's disease or a disease or condition involving aberrant apoptosis. They are also useful in research and diagnostics for modulating the expression of amyloid beta protein precursor. The present sequence is human amyloid beta precursor DNA amplifying PCR primer.
                                                                                                       New RNA molecule less than 49 bases and having a mediate degradation or block translation of mRNA transcriptional product of a target gene, useful disease or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                 21-AUG-2002; 2002US-0405193P
03-SEP-2002; 2002US-0408152P
20-MAY-2003; 2003US-0472387P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RNA interference; RNAi; short interfering RNA; siRNA; human; insulin-like growth factor binding protein-5; cytostatic; neuroprotective; nootropic; gene silencing; IGFBP-5; DNA-RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 9 A; 7 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                              WPI; 2004-226852/21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-MAR-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA1 for human insulin-like growth factor binding protein-5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-MAY-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADL70482 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                treating Alzheimer's disease or a disease
                                                                                                                                                                                                                                                                                                                                                                                 (UYBR-) UNIV BRITISH COLUMBIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5170 CACAGGGTGAGCCCCAAAA 5188
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                                                  SEQ ID NO 27, 63pp;
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                                                                                                                                                                                                                                                                                                                         Gleave
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nilarity 89.5%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "OTHER= dTdT"
                                                                                                                                                                                                                                                                                                                            Signaevsky
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.8; DB 1;
Pred. No. 7.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                            Į,
                                                                                                                                                                                                                                                                                                                               Beraldi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   involving aberrant apoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IGFBP-5; DNA-RNA hybrid; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ϊ,
                                                                                                                                                                                                                                                                                                                               M
                                                                                                                                   sequence effective to
that is the
for treating Alzheimen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 21;
                                                                                                                                                                                                                                                                                                                            Trougakos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                         Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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Claim 5;

present sequence is the sense strand

of a short interfering RNA

English

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RESULT 629
ADL60643/c
ID ADL606
XX ADL606
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XX ADL606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Sim
Matches 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (siRNA) targeted to nucleotides 1225-1243 of human insulin-like growth factor binding protein-5 (IGFBP-5) cDNA. The antisense strand is also provided ADL70483. The siRNA can be used to interfere with the expression of IGFBP-5. Inhibition of IGFBP-5 expression can delay the progression of hormone-regulated (prostatic or breast) tumour cells to hormone independence, provide a method for the treatment of hormone-regulated cancers, and inhibit or delay the growth and metastatic progression of prostate, breast and other IGF-1 sensitive tumours in bone. siRNAs of the invention can be used alone or in combination with other chemotherapy or sarcomas such as osteosarcoma, renal cell carcinoma, breast cancer, bladder cancer, lung cancer, colon cancer, ovarian cancer, anaplastic large cell lymphoma and melanoma, and also for the treatment of
                                                                                                                              New nucleic acid encoding human OATP2 or cMOAT protein, useful diagnosing, treating or preventing diseases or disorders, e.g. inflammatory, cardiovascular, hyperproliferative, neurological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene therapy inflammatory
                                                                                                                                                                                                                                                                          (TSUC/)
                                                                                                                                                                                                                                                                                                                           21-SEP-2001; 2001US-0324172P
27-NOV-2001; 2001US-0333700P
                                                                                                                                                                                                                                                                                                                                                                         20-SEP-2002; 2002US-00252155
                                                                                                                                                                                                                                                                                                                                                                                                                                         US2004068096-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     coronary heart disease; cancer; wound healing; ds; SNP; single nucleotide polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    gene therapy; human; OATP2; cMOAT; hepatic disease; metabolic disinflammatory disease; cardiovascular disease; hyperproliferative neurological disease; infectious disease; liver disease; high cholesterol; hypertension; congestive heart failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human organic anion transport protein SNP region #369.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADL60643 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                               Tsuchihashi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Alzheimer's
                                                                                                                 infectious
                                                                                                                                                                                                                                                             (KIRC/)
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                                                                                                                                                                                               2004-304621/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens
                                                                                                                                                                                                                                                                            TSUCHIHASHI Z.
HUI L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 BP; 6 A; 6 C; 5 G; 2 T; 2 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGCUGACCCAGUCCAAGTT 21
                                                                                                                   diseases.
                                                                                                                                                                                                                       27
                                                                                 SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                             Hui
                                                                                 ID NO
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78.9%;
                                                                                419; 296pp; English
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                                                                                                                                                                                                                            Kirchgessner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2;
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No. 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease;
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The invention relates to an isolated nucleic acid derived from a hugene encoding a protein, i.e. human OATP2 protein or human CMOAT puwhere the nucleic acid comprises at least one polymorphic position.

encoded protein,

composition

protein,

Sequence

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BP; 1 A;

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0 SNP

ä 0 region.

organic

anion

transport 8

protein 11 Ŧ;

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RESULT 630
ADL60642/c
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Best Local Similarity
Matches 17; Conserv
                                                                          The invention relates to an isolated nucleic acid derived from a human gene encoding a protein, i.e. human OATP2 protein or human CMOAT protein, where the nucleic acid comprises at least one polymorphic position The nucleic acid and the encoded protein, kits and composition are useful in diagnosing, treating or preventing diseases or disorders, e.g. hepatic, metabolic, inflammatory, cardiovascular, hyperproliferative, neurological, infectious diseases, liver disease, high cholesterol, hypertension, congestive heart failure or coronary heart disease and cancer and promotes wound healing. The present sequence represents the a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             diagnosing, treating or preventing diseases or disorders, e.g. hepatic, metabolic, inflammatory, cardiovascular, hyperproliferative, neurological, infectious diseases, liver disease, high cholesterol, hypertension, congestive heart failure or coronary heart disease and cancer and promotes wound healing. The present sequence represents the human organic anion transport protein SNP region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 1 A; 8 C; 1 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                      New nucleic acid encoding human OATP2 or cMOAT protein, useful diagnosing, treating or preventing diseases or disorders, e.g. inflammatory, cardiovascular, hyperproliferative, neurological infectious diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tsuchihashi Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene therapy; human; OATP2; cMOAT; hepatic disease; metabolic disease; inflammatory disease; cardiovascular disease; hyperproliferative diseasenurological disease; infectious disease; liver disease; high cholesterol; hypertension; congestive heart failure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human organic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUL-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-SEP-2001; 2001US-0324172P
27-NOV-2001; 2001US-0333700P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-APR-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      coronary heart
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (KIRC/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (TSUC/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2004-304621/28.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ) TSUCHIHASHI Z.
) HUI L.
) KIRCHGESSNER T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GAAAGCAGAGAAAGAGGGA
                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           anion transport protein SNP region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   disease; cancer; wound healing; ds; SNP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hui L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        polymorphism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
                                                                                                                                                                                                                                                                                                                                                                                418; 296pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kirchgessner T;
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. 7.8e+02;
2;
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RESULT 631
AAT92356/c
ID PART9233
XX AAT9233
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XX AAT923
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Best Local Similarity
Matches 17; Conserv
                     Achiral linker reagents have been developed for the incorporation of multiple amino groups into oligonucleotide. The present sequence represents a modified oligodeoxyribonucleotide. The achiral linker reagents can be used for incorporation of multiple primary amino groups or reporter groups into oligonucleotides. They are compatible with conventional DNA synthesis following the bhosphoramidite methodology, an can be incorporated in good yields. The linker reagents may be used for labelling of oligonucleotides. They may also be used for use as primers in the polymerase chain reaction or in nucleic acid sequencing reactions, for production of affinity matrices for detection of the biomolecules, for production of affinity matrices for detection of nucleic acid sequences, for cloning recombinan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Amino modified oligodeoxyribonucleotide; oligonucleotide; achiral linker reagent; 5 (aminomethyl)-1,3 benzenedimethanol; N-fluresceinyl-(5-aminomethyl)-1,3-benzenedimethanol; hybridisation probe; PCR primer; nucleic acid sequencing; hybridisation grobe; PCR primer; nucleic acid sequencing; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New achiral linker reagents - useful
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Pred. No. 7
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                                                                                                      This exon specific primer used in the PCR amplification of a short content associated sequence. The gene region corresponding to short stature has been identified as a region of approximately 500 kb in the CR ARI region of the X and Y chromosomes. Three genes in this region have been identified as candidates for the short stature gene. These genes CR were designated SHOX (also referred to as SHOX3) or HOX93), pET92 and CR SHOX (SHOX TIME homeobox gene on chromosome three). The SHOX gene has two separate splicing sites resulting in two variations SHOXa and SHOXb. The specification provides sequences of SHOX (short stature homeobox—CR specification than ANJ35610 to ANV35621 and protein sequences of the SHOX genes as shown in ANV35610 to ANV35621 and protein sequences of the human growth protein transcription factor SHOXA, SHOXb and SHOT as shown and SHOT as shown can be used to develop agents for the treatment of short stature contains a shown of the short stature can be used to develop agents for the treatment of short stature contains a shown and store human growth disorders. The products can also be used for the sequence in the sequence in sequences and diseases involved with disturbance in the sequence in sequences and diseases involved with disturbance in the sequence in the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New human growth genes - used to develop products for the diagnosis and treatment of human growth defects such as short stature, e.g. Turner's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1998-271719/24
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16-JAN-1997;
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                                                                          such as osteoporosis and diseases calcium regulation
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Matches 17; Conservative
                                                                                                                                                                                                                                                                                     or upregulation of telomerase expression in cancer. Substances, identified in the methods, can be used to block transcription from the TR gene promoter through interaction of the 5' regulatory sequences. These substances, e.g. antisense oligonucleotides, transcription factors, peptide nucleic acids and factors that disrupt signal transduction, are useful for cancer therapy. In particular, gene therapy vectors (especially pGTG2-codhupp) comprising the promoter and a viral thymidine kinase gene can be used to convert a prodrug, e.g. gancyclovir, so that neoplasia can be controlled or treated. Direct down-regulation of telomerase RNA gene through manipulation of transcription factors may be effective anticancer therapy and the cloning of the hTR gene promoter allows the analysis of therapy and the cloning of the hTR gene promoter activity. Sequences AAZ07696-321 represent PCR primers used in cloning and mutagenisis of human TR gene (hTR) promoter region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene therapy; thymidine kina mutagenesis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to promoter regions from mouse and human telomerase RNA (TR) component genes. The TR gene promoter can be linked to a heterologous gene, especially a gene encoding a cytotoxin, for therapy of cancer, especially neoplasias. The telomerase is necessary for the unrestricted proliferative capacity of many human cancers. Mutation or dysregulation of the telomerase repression pathway may cause reactivation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Telomerase RNA; TR; promoter; cytotoxin; cancer; neoplasia; hTR; gene therapy; thymidine kinase gene; anticancer therapy; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ07304 standard; DNA; 22 BP.
                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Fig 12; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mouse and human telomerase RNA gene promoters, useful for tumor specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1999-479183/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-AUG-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (CANC-) CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3988
2179 CATTACCTTGCCCAGGCTC 2197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sapiens.
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                                                                                          0.3%;
l Similarity 89.5%;
17; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GCTGAGCCTGGAGCTGTGG 4006
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                                                                                                                                                                                                                                   22
                                                                                                                                                                                                                            BP; 4 A; 4 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
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                                                                                                0;
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0; Mismatches 2;
                                                                                                                              Score 15.8; DB 1; Length 22; Pred. No. 7.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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                                                                                            Query Match
Best Local Similarity
                                                                          Matches
                                                                                                                                                                                   With protein EF1 gamma. Post-translational phosphorylation and dephosphorylation modulates the effect of the FEZ1 protein. Inhibitors of FEZ1 gene expression are useful for inducing cells to proliferate. Compounds which modulate FEZ1 association with tubulin are useful for alleviating tubulin hyper- or hypo- polymerisation disorders, such as those associated with aberrant initiation of mitosis, modulation of the initiation and rate of cell proliferation and cell growth, modulation of cell shape, cell rigidity, cell motility, rate and stage of cellular DNA replication, intracellular distribution of organelles, metastatic potential of cell and cellular transformation from a non-cancerous to cancerous phenotype. Compounds which modulate FEZ1 binding and phosphorylation are also useful for alleviating a disorder, such as tumorigenesis, tumour survival, growth and metastasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; FBZ1 gene; tumour suppressor gene; 8p22; cancer; tumour growth; tumour proliferation; tubulin; microtubule; protein EF1-gamma; tubulin polymerisation disorder; mitosis initiation; cell proliferation; cell growth; cell shape; cell rigidity; cell motility; DNA replication; tumorigenesis; tumour survival; metastasis; PCR primer; ss.
                                                                                                                                                    Sequence 22 BP; 2 A; 10 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                          PCR primers AAA64531-32 were used to amplify a fragment of the human FE: gene. FEZ1 is a tumour suppressor gene, located at chromosome location 8p22. Decreased or no expression of FEZ1 is detected in a variety of cancer cells. Expression of FEZ1 inhibits tumour growth and proliferation. FEZ1 also interacts with tubulin, with microtubules, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Croce CM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polynucleotide homologous with a portion of one strand of the human FEZ1 gene, useful for alleviating abnormal cell proliferation such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA64532 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 1; Page 45; 255pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYJE-) UNIV JEFFERSON THOMAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-FEB-1999; 99US-0121537P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-FEB-2000; 2000WO-US004950.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PCR primer G2 used to amplify exon 2 of human FBZ1 gene
                                    2640 CCTGCAGCTGCTGCAG 2658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-558396/51
4
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                                                                          Conservative
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                                                                                            89.5%;
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                                                                          0;
                                                                                            Score 15.8; DB 1;
Pred. No. 7.8e+02;
                                                                            Mismatches
                                                                                                                DB 1; Length 22
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                                                                            Gaps
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RESULT 635 AAF83320/c

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RESULT 636
ABL45282
ID ABL452
XX ABL452
XX 11-APR
XX 11-APR
XX Human;
XX Human;
XW PCR px
XX Homo 8
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to SAPL [SIT4-(sporulation-induced transcript4) associated proteins-like] polypeptide, selected from SAPLa polypeptide isoforms and SAPLa bolypeptide isoforms. The SAPL polypucleotides are useful in gene therapy for treating and preventing insulin-dependent diabetes mellitus (IDDM). Fragments of the SAPL DNA are useful as primers and probes. The SAPL polypeptides are useful in screening for a substance e.g., a peptide or chemical compound, which interacts and/or binds with them. Sequences AAPB3318-350 represent PCR primers specific for the SAPL DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isoforms of novel gene arising from alternative splicing and encoding highly related proteins termed as SAPLa and SAPLb, from the IDDM4 locus on human chromosome 11q13, useful for treating IDDM and other diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SAPL; SIT4; SIT4 associated proteins like; human; antidiabetic; sporulation-induced transcript 4; SAPLa; SAPLb; gene therapy; IDDM; insulin-dependent diabetes mellitus; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 22 BP; 7 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 14; Page 98; 129pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-300338/31.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        09-JUL-2001
   JP2001321190-A
                              Homo sapiens.
                                                                        Human; chromosome 1p36-35; chromosome 21q22.1; genetic analysis; genome;
                                                                                                    Human chromosome 1p36-35 PCR primer SEQ ID NO:2326.
                                                                                                                                 11-APR-2002
                                                                                                                                                              ABL45282;
                                                                                                                                                                                         ABL45282 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Todd JA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (WELL ) WELLCOME TRUST LTD. (MERI ) MERCK & CO INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAF83320 standard; DNA; 22 BP
                                                         primer; ss.
                                                                                                                                                                                                                                                                                            2359 CACCCCATCCCTCTGAGCT 2377
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                                                                                                                                                                                                                                                                                                                                      Similarity 89.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Twells RCJ,
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                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                 3 C; 8 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                    0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            невв JW,
                                                                                                                                                                                                                                                                                                                  0,
                                                                                                                                                                                                                                                                                                                                                    Score 15.8; DB 1;
                                                                                                                                                                                                                                                                                                                                      Pred.
                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                        No.
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                                                                                                                                                                                                                                                                                                                                                    Length 22;
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RASULT 637
ABKS2714
ID ABKS277
XX ABKS27
XX ABKS27
XX Human
XX Human
XX PCR; p
XX PCR; p
XX JP2002
XX JP2002
XX JP2002
XX JP2002
XX JP3007
XX GN3-OCT
XX ON-OCT
XX (KBIO-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention describes a method of arraying genome clones. The method comprises; (a) clones of the genomic libraries contained in clones in the genomic libraries contained in multiwell plates numbered for discrimination are mixed in each of the multiwell plates; (b) a primer designed based on the chromosome marker sequence is added to the mixture to carry out an amplification reaction; (c) a signal corresponding to the marker is detected from the resultant amplified product to specify the discrimination Nos. of the multiwell confide the markers is changed so that the same discrimination Nos succeed to the maximum in the specified discrimination Nos. to array the multiwell plates; (e) the clones in the multiwell plates of the specified confideration Nos. are mixed respectively in each wells of longitudinal cardinariation Nos. are mixed respectively in each wells of longitudinal cardinariation Nos. are mixed respectively in each wells of longitudinal cardinariation the amplified by using the above primer; (g) signals are detected from the amplified products; (h) the clones in the multiwell plates are specified from the detected result; and (i) the clones are reconstituted as the positions on the chromosome and arrayed. The microarray is useful for gene analysis. ABL42957 to ABL45323 to ABL45634 represent PCR primers for human chromosome 21q22.1, which are generally claimed for use in the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local !
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                       03-OCT-2000; 2000JP-00304143.
                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                                                                                        Human bladder cancer antigen KU-BL-1, PCR primer #10.
                                                                                                                                                                                                                                                                                                             27-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                   ABK52714 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22 BP; 5 A; 12 C; 0 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Page 50; 528pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Arraying genome clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-MAR-2000; 2000JP-00066716
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                                                                                                             16-APR-2002.
                                                                                                                                               JP2002112779-A
                                                                                                                                                                                                                                      Human; bladder cancer antigen KU-BL-1; immunotherapy; cytostatic; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (RIKA) RIKAGAKU KENKYUSHO.
(GENO-) GENOTEX YG.
                                                                                                                                                                                                                     primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                343 CTACCACTCCCCCTCTATC 361
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.8;
Pred. No. 7.
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(KBIO-) GH KBIO GIJUKU.

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RESULT 638
ABV99963/c
ID ABV999
XX ABV999
XX 28-MAR
XX Human
XX G prot
KW G prot
KW hypote
KW manic
KW manic
KW manic
KW manic
KW FR2823
XX FR2823
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The present invention relates to a human G protein coupled receptor and its coding sequence. The coding sequence is useful for identifying specific modulators. In addition, the coding sequence, and related sequences, also host cells containing them, are useful for screening for agonists and antagonists of the G protein coupled receptor. The coding sequence is also useful for recombinant production of the G protein coupled receptor, and as source of amplification probes and primers for identifying pathological alleles or species homologs. Measuring the expression of the G protein coupled receptor and its coding sequence are useful for diagnosis, prognosis and monitoring of diseases associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  manic depression; delirium; dementia; epilepsy; migr
circadian rhythm disorder; cognitive impairment; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G protein;
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                                                                                                                                                                                                                                                                                                                                                                     diabetes.
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The invention relates to an isolated nucleic acid (N1) comprising at cleast 20 but not more than 1500 consecutive nucleotides of the optineurin promoter appearing as ADE13890. Also included are the optineurin promoter coperably linked to a heterologous nucleic acid, a nucleic acid capable of detecting a single nucleotide polymorphism (SNP) in the optineurin promoter promoter, a host cell comprising the promoter operably linked to a comprising of the promoter operably linked to a comprising detecting a polymorphism of a promoter region of the optineurin gene, associated with a glaucoma composition of the optineurin promoter sequence variation in a sample containing DNA, detecting a SNP sequence variation in a sample containing DNA, determining the presence of an optineurin promoter sequence variation in a sample containing DNA, determining the presence or increased susceptibility to glaucoma or to a progressive ocular hypertensive of disorder resulting in loss of visual field in a patient (or the severity or progression of glaucoma in a patient, comprising providing amplification reaction primers that direct amplification of a selected
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid sequences of the optineurin gene are useful to detect polymorphisms particularly single nucleotide polymorphisms in the optineurin promoter to diagnose, prognose and treat glaucoma and related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 11;
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                                        The invention relates to diagnosing a predisposition to fat deposition leanness in a subject comprising detecting the presence or absence of polymorphic variation associated with fat deposition at a polymorphic site in a PLAZGIB nucleotide sequence in a nucleic acid sample from a subject, where the presence of the polymorphic variation indicates a predisposition to fat deposition in the subject. The polymorphic variation is a guanine at position 7328 or thymine at position 9182 of the present sequence. The method is useful for diagnosing a predisposition to fat deposition or leanness in a subject, and consequently for diagnosing a predisposition to non-insulin dependent diabetes mellitus (NIDDM) in a subject and conditions such as hyperinsulinemia, hypertension, glucose intolerance, dyslipidemia, hypercospulability, or microalbuminuria, which can lead to early
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid region containing the variation within the optineurin promoter and amplifying the DNA) and detecting a polymorphism (comprising obtaining a sample containing human genomic DNA, providing a nucleic acid capable of detecting a SNP located within an optineurin promoter, and detecting the polymorphism). The invention is used to diagnose and prognose glaucoma and also to treat glaucoma related disorders. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Diagnosing a predisposition to fat deposition or leanness, useful for diagnosing a predisposition to e.g. diabetes or hypertension, comprises detecting the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the PLA2GIB nucleic acid from the presence of a polymorphism in the presence
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                                               This invention relates to a novel candidate therapeutic agent useful for CC fat reduction and disorders related to fat depositions. Specifically, it CC refers to polymorphic variations in the phospholipase A2 (PLAZGIB) DNA, CC which is located on chromosome 12q24 and has been associated with central fat deposition. The present invention describes methods to detect the presence or absence of these single nucleotide polymorphisms of PLAZGIB, CC in particular G7328A and T9182G, and subsequently provide treatment that creduces fat deposition. This treatment may consist of an appetite compressant, a lipase inhibitor, a phospholipase inhibitor, an exercise cregimen, a distary regimen, psychological counselling, psychotherapy or a psychotherapeutic. Accordingly, PLAZGIB is a target for reducing fat deposition and it can be used to treat both obesity and non-insulin consolidate mellitus (NIDDM), as well as cardiovascular disorders coligonucleotide sequence is a PCR primer used to amplify fat sand rat plazgib DNA, used in an exemplification of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 5; Page 80; 116pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Identifying a candidate therapeutic for fat reduction, useful treating diabetes, by introducing a test molecule to a system PLA2G1B protein or nucleic acid, and determining the presence
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No. 7.8e+02
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CC pectoris, cardiac infarct, coronary spasm, cardiac insufficiency, stroke C and peripheral occlusive disease); chronic inflammatory and autoimmune C diseases (rheumatoid arthritis, psoriatic arthritis, Crohn's disease); CC diabetes types I and II (and complications such as nephropathy); multiple CC sclerosis; sarcoidosis, collagenosis and vasculitis, including CC glomerulonephritis; acute and chronic rejection reactions of transplanted CC organs; graft versus host disease and ischaemic/reperfusion injury during CC surgery; allergic contact dermatitis, (pre-)eclampsia; hypertension and CC thickening); chronic kidney disease; chronic obstructive lung disease; CC tubercular pericarditis, Lyme borreliosis and its consequences); CC tubercular pericarditis, Lyme borreliosis and its consequences, particularly cortal hypertension and fibrosis, also opportunistic infections, CC particularly Pneumocystis carinii pneumonia. Also, detecting the -786C/T polymorphism in the eNOS gene is useful for diagnosis. The present CC sequence represents a double-stranded oligonucleotide of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New double-stranded oligonucleotides, useful for treating e.g. cardiovascular disease and arthritis, are decoys that lift inhibition
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Crohn's disease; diabetes; nephropathy; multiple sclerosis; sarcoidosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to double-stranded DNA oligonucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Double stranded oligonucleotide #3
      tubercular pericarditis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (AVON-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12-SEP-2002; 2002DE-01042319.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-SEP-2002; 2002DE-01042319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4109 AGCCAGCCAGGGTGGAGCT 4127
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AVONTEC GMBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AGCCAGCCAGGGAAGAGCT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hecker
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  36pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          to double-stranded
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      German
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 7.00
0; Mismatches
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Pred. No. 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             are decoys that lift inhibition
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            u; o
                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA oligonucleotides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .8e+02;
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-786C/T

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RESULT 644
ABZ07924/c
ID ABZ079
XX ABZ079
XX ABZ079
XX ABZ079
XX ABZ079
XX PT; le
XW Stherc
XX FT; le
XW SS.
WO200;
XX SS.
PD 25-JUJ
XX 25-JUJ
XX 25-JUJ
XX WO19
PD 25-JUJ
XX 25-JUJ
XX 25-JUJ
XX 25-JUJ
XX 26-OC
PR 08-JUJ
XX WO19
PF 22-OC
XX WO19
PF 22-OC
XX WO19
PF 22-OC
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Query Match
Best Local Similarity
Matches 26; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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Best Local (
                                                                                                                                                                                  The invention relates to a system for detecting gene expression, which comprises one or two isolated DNA molecules that detect expression of a gene, where the gene corresponds to any of 8143 oligonucleotides (ABZ00010-ABZ08152) each having 50 base pairs (bp). The system is useful for leukocyte expression profiling. It is particularly useful for diagnosing a disease, monitoring (rate of) progression of a disease, predicting therapeutic outcome, determining prognosis for a patient, predicting disease complications in an individual or monitoring response to treatment in an individual. The diseases include cardiac allograft rejection, kidney allograft rejection, liver allograft rejection, therapeutic outcome heart failure, systemic lupus erythematosus, congestive heart failure, systemic lupus erythematosus,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human leukocyte
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                portal hypertension and fibrosis, also opportunistic infections, particularly Pneumocystis carinii pneumonia. Also, detecting the -786C/polymorphism in the eNOS gene is useful for diagnosis. The present sequence represents a double-stranded oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New system
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wohlgemuth J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-OCT-2000; 2000US-0241994P.
08-JUN-2001; 2001US-0296764P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             T7; leukocyte; gene expression profiling; allograft rejection; atherosclerosis; congestive heart failure; systemic lupus eryt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-JAN-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABZ07924;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABZ07924 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              22-OCT-2001; 2001WO-US047856.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rheumatoid arthritis; osteoarthritis; cytomegalovirus; infection; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22 BP; 2 A; 8 C; 6 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (BIOC-) BIOCARDIA INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    congestive heart failure,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4109 AGCCAGCCAGGGTGGAGCT 4127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1; Page 583; Opp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19
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                                                                                                                                                              arthritis, osteoarthritis or cytomegalovirus infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    for leukocyte expression profiling, diagnosing a disease, or (the rate of) progression of a disease, e.g. atherosclerosis ive heart failure, comprises diagnostic oligonucleotides.
  Conservative
                                                                                                             BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                                                                             A; 9 C; 18 G;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.3%;
                         0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matcuk G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  50
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  0
  Score 15.8; DB 1;
Pred. No. 7.9e+02;
0; Mismatches 17
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Pred. No. 7.
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                                                                                                             11 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Altman P, P, Johnson F;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8; DB 1;
7.8e+02;
                                                                                                                U, 0 Other;
                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prentice J,
                                                      Length 50;
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     Gaps
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RESULT 645
ADP10317/c
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XX ADP103
XX ADP103
XX Transp
XX transp
XX transp
XX Homo a
XX W02004
XX W02006
XX W02006
XX W01;
Z1-MAY
XX 24-APF
PR 20-DEC
XX W01;
Z1-MAY
XX CEXPR-
XX W01;
Z1-MAY
XX Transp
PF 24-APF
PR 20-DEC
XX WPI;
Z1-MAY
XX CEXPR-
XX WPI;
Z2 W011g
Z2 W011g
Z3 W011g
Z4 W01;
Z5 Claim
XX The pr
CC reject
CC Transp
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                                                                                                                Matches
                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                           rejection, e.g. cardiac or kidney transplant rejection, in an individual comprises detecting the expression level of one or more genes. The methods, system and kits are useful in diagnosing or montcoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual. The method is also useful in assessing the immune status of an individual. The methods are also useful in diagnosing and monitoring diseases that involve the immune system, e.g. rheumatoid arthritis, lupus, inflammatory bowel diseases, multiple sclerosis, HIVAIDS or viral, bacterial or fungal infection. The present sequence represents a 50 mer oligonucleotide marker for diagnosis and monitoring of allograft rejection and other disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diagnosing or monitoring transplant rejection, e.g. heart, kidney, liver, pancreas, pancreatic islet, lung, bone marrow or stem cell transplant rejection, in an individual, comprises detecting the expression level of
                                                                                                                                                                                                                              Sequence 50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention relates to diagnosing or monitoring transplant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the genes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wohlgemuth J, Rosenberg S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-APR-2002; 2002US-00131831.
20-DEC-2002; 2002US-00325899.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-APR-2003; 2003WO-US012946
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transplant rejection; immune system; rheumatoid arthritis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            50-mer oligonucleotide marker probe of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-AUG-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADP10317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADP10317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2004042346-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          inflammatory bowel disease; multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EXPR-) EXPRESSION DIAGNOSTICS INC.
                                                                                                                                         Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4022 CTCACTITGTGGCTCTCCAAGGGGGCCATGTGGACACATCCCT 4064
                                                   4022 CTCACTTTGTGGCTCTCCAAGGGGGCCATGTGGACACATCCCT 4064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2004-400724/37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               50
                                                                                                             26;
                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 326; 1762pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ĊŢĊĄĊĊĊŢĠŢĠĄĊĄĊĄĠĊĊĄĠĄĠĠĊĄĄĄŢĠGŢŢĊĊĄŢĠŢĊĊĊŢ
                                                                                                                                                                                                                                 BP; 12 A; 9 C; 18 G; 11 T; 0 U; 0 Other;
                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fry K,
                                                                                                                                     0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Woodward
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                50
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                뫄
                                                                                                             0;
                                                                                                                                            Score
Pred.
                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₽,
                                                                                                                                         15.8; DB 1;
No. 7.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ę
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sclerosis; HIV; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ,z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prentice
                                                                                                                                                                    Length 50;
                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               œ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      #326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Morris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       kidney, liver,
                                                                                                                0;
                                                                                                                Gaps
                                                                                                                   0,
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RESULT 646
AAT27910/c
ID AAT27910 standard; DNA; 20

BP.

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RESULT 647
AAQ37956/c
ID AAQ37956 standard; DNA; 22 BI
XX
AC AAQ37956;
XX
DT 25-MAR-2003 (revised)
DT 11-JUL-1993 (first entry)
XX
DE Sequence of transcription factor: YV1; mc
XX
Transcription factor: YV1; mc
XX
Epstein-Barr virus; oncogene;
XX
OS Homo sapiens.
                                                                                                                                                                                                                                                                                                  문
                                                                                                                                                                                                                                                                                                                         S
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect cpd. simple sequence repeat (SSR), and an adaptor directed primer, comprising a sequence complementary to an adaptor directed primer, comprising a sequence of a SSR primer, which is flanked at its 5'-end by degenerate nucleotides. The method represents a modified amplified fragment length polymorphism assay, which is partic.
                                                                                   Sequence of transcription factor position (P5+1) mt2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; 5'-anchored; ss.
                                   Epstein-Barr virus; oncogene;
                                                   Transcription factor: YY1; modulatory
                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 0 A; 8 C; 0 G; 9 T; 0 U; 3 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modified amplified fragment length polymorphism assay - for detection polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Morgante M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9617082-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 76; 173pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-277795/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5'-anchored simple sequence repeat primer DVD(TC)8.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              28-JAN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAT27910;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (DUPO ) DU PONT DE NEMOURS & CO E I.
                                                                                                                                                                                                                                                                                                                              1182 AGAAAGAGAGAGAGAA 1199
                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                AGAGAGAGAGAGAGAH 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                              comparisons
                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vogel JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94US-00346456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95WO-US015150
                                                                                                                                                                                                                                                                                                                                                                             88.9%;
                                                                                                                                                                                                                                                                                                                                                                                              0.3%;
                                                                                                                                                                                                                   ₽P
                                                                                                                                                                                                                                                                                                                                                              ۲,
                                                                                                                                                                                                                                                                                                                                                                             Score 15.6; DB 1.
Pred. No. 8.2e+02
                                      88
                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                    (TF)
                                                     adeno-associated virus
                                                                                                 YY1 mutant binding sequence at
                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                    P5+1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         œ,
                                                                                                                                                                                                                                                                                                                                                              0
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13-NOV-1992;

92US-00977284. 93WO-US010964.

12-NOV-1993; 26-MAY-1994 Synthetic

Type II procollagen; COL2A1; amplification; primer; polymerase chain reaction; PCR; osteoarthritis; car

cartilage;

II procollagen PCR primer 83.

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RESULT 648
PAQ65892
ID AAQ658
XX AAQ658
XX 25-MAR
DT 25-MAR
DT 22-DEC
XX Type I
XX Type I
XX Polyme
XX Polyme
XX W0941]
XX W0941]
XX PF 12-NO1
XX PF 12-NO1
XX PF 11-NO1
XX PF 11-NO1
XX PR 13-NO1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The cDNA of transcription factor (TF) YY1 (AAQ37948) was derived from CHeLa cells derived from cervical carcinoma from clone p14-1 or pYY1 of the D98/AH-2 library. TF YY1 has a mol. wt. of 68 kD and is capable of binding to a sequence between -50 and -70 (P5-60 site) and to the transcription initiation region of the promoter (P6+1 site) of an adenosate accided virus (AAV). It represses transcription directed by a TATA celement plus initiator (Inr) sequence. It can mediate transcription through an Inr sequence or element. The initation site is of a lymphocyte specific terminal deoxymuclectidyltransferase gene (TdtInr) or human leucocyte IFN gene (LeIF-J Inr). YY1 can effect latency of viruses. It is a zinc finger protein and has a glycine rich sequence, as well as committed twestion of the two sequences (see AAQ37950-52, AAQ37954-56) were placed directly upstream of the TATA box in construct pTI to the state the functions of these binding sites within transfected cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Purified and isolated mammalian transcription factor YY1 - used to modulate transcription in adeno-associated virus type 2- Epstein-Barr viruses and oncogenes; useful in diagnosis and screening.
                                                                                                                                                                                                                                                                                                                                           AAQ65892 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22 BP; 4 A; 4 C; 10 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                              25-MAR-2003
22-DEC-1994
                                                                                                                                                                                                                                                                                                             AAQ65892;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1993-093929/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Shi Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-AUG-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-MAR-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W09304076-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYPR-) UNIV PRINCETON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-AUG-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Updated on
                                                                                                                                                                                                                                                                                                                                                                                                                                                       3026 CTCGCTGCCTCCTGGAGACCCT 3047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18;
                                                                                                                                                                                                                                                                                                                                                                                                                           22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Seto E, Shenk T;
                                                                                                                                                                                                                                                                                                                                                                                                                           CCCGCTTCCAACTGGAGACCCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fig 5; 100pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                            (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91US-00746485.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          92WO-US006840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
                                                                                                                                                                                                                                                                                                                                            ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.6; DB 1; Length 22; Pred. No. 8.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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(UYJE-) UNIV

밀

Ala-Kokko L, JEFFERSON THOMAS

Williams

, 5

Ritvaniemi P,

Baldwin

ņ

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RESULT 649
AAV02438
ID AAV0244
XX AAV024
XX O6-APR
XX Human
XX G prot
KW pituit
KW pituit
KW therap
XX Homo s
XX Hom
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  δ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                     15-MAR-1996;
12-AUG-1996;
18-SEP-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        G protein-coupled receptor; ligand binding; pharmaceutical; modulator; pituitary; central nervous system; pancreas; prophylactic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 18 claims primers for use in detecting mutations in a mammalian gene for a structural protein of cartilage comprising a sequence identified in Table I (Page 18-31). Table I includes 179 primer sequences (see AAQ65728-Q65906). The following details are given for primer 83: Alt. code: DH-75 Region/exon: 49 Direction: sense Primer position: 20047 (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV02438 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 22 BP; 8 A; 2 C; 7 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Detecting genetic pre-disposition to osteoarthritis - and other diseases involving mutation in cartilage protein genes, by amplification and analysis of DNA and comparison with standards.
                                                                         Example 31; Page 133; 258pp;
                                                                                                                         Ligand peptide for G function in the centr
                                                                                                                                                                                                                                                                                                                                                                                                                                               28-DEC-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          W09724436-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    06-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAV02438
                                                                                                                                                                                                   WPI; 1997-363672/33
                                                                                                                                                                                                                                                                                                                       (TAKE ) TAKEDA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUL-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 therapeutic agent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          iomo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2755 GTAGAAACAGACATGGAGCTCT 2776
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18;
                                                                                                                                                                                                                                                   ပ်စ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GAAGAATAGACATGGTGCTGT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 30; 112pp;
                                                                                                                                                                                                                                                                          Habata Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                     95JP-00343371.
96JP-00059419.
96JP-00211805.
96JP-00246573.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ahmad
                                                                                                                                                                                                                                                                                                                          CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96WO-JP003821.
                                                                                                                         for G protein-coupled receptor central nervous system, pancre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Z
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              81.8%;
                                                                                                                                                                                                                                                                          Kawamata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 8.36
0; Mismatches
                                                                         English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PCR primer
                                                                                                                                                                                                                                                                       ۲
                                                                                                                                                                                                                                                                       Новоуа
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8.3e+02
                                                                                                                         pancreas
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this ligand may be used as a pituitary function modulator, a central nervous system modulator or a pancreatic function modulator. This ligand could have specific applications as a prophylactic or therapeutic agent for dementia, depression, hyperkinetic syndrome, disturbance of consciousness, anxiety syndrome, schizophrenia, trauma, growth hormone secretory disease, hyper- and polyphagia, hypercholesterolaemia, hyperglyceridaemia, hyperlipidaemia, hyperprolactinaemia, diabetes, cancer, pancreatitis, renal disease, Turner's syndrome, neurosis, asthma, rheumatoid arthritis, spinal injury, transient brain ischaemia, epilepsy, amylotrophic lateral sclerosis, acute myocardial infarction, infertility, spinocerebellar degeneration, bone fracture, trauma, atopic dermatitis, osteoporosis and/or oligogalactia. Assays can also be developed to screen compounds which are capable of altering the binding activity of the
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein-coupled receptor protein. Pharmaceutical compositions containing
22 BP; 1 A; 6 C; 7 G; 8 T; 0 U; 0 Other;
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Query Match
Best Local S
Matches 18
Human mutant KCNQ2 screening
                  22-JUL-1999
                                                          AAX57086 standard; DNA;
                                                                                                                              3260 ACCTGGCCTCTGTGCTTAGTGC 3281
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                   (first entry)
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primer 13.
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KCNQ2; KCNQ3; human; murine; potassium channel; diagnosis; benign familial neonatal epilepsy; BFNE; juvenile myotonic rolandic epilepsy; mutant; treatment; screening; epilepsy; gene therapy; drug screening; primer; ss. Synthetic. JME;

06-MAY-1999 W09921875-A1 sapiens

(UTAH) UNIV UTAH RES FOUND

24-OCT-1997; 23-OCT-1998;

97US-0063147P 98WO-US022375

Ä, Leppert MF, Charlier C;

1999-312938/26

Nucleic acid encoding potassium channels KCNQ2

Claim Page 144; 195pp; English.

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AR This invention describes novel human and mouse potassium channel proteins KCNQ2 and KCNQ3. Detecting mutations in sequences that encode KCNQ2 or KCNQ3, or the loss of one copy of these genes, is used for diagnosis and prognosis of benign familial neonatal epilepsy (BFNR), juvenile myotonic epilepsy (JME) or rolandic epilepsy (RE). Cells (or transgenic animals) that express wild-type or mutant KCNQ2 or 3 (also the proteins themselves in cell-free form) are used to screen for agents that can be used to treat or prevent these forms of epilepsy. Fragments of the encoding nucleic acids are used as probes or primers, either for detecting

PCR primers type ligand

AAV02438-V02440 are used to amplify a novel bioactive human cDNA used in an assay to monitor ligand binding to the G

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ANOSITY 651
ANOSITY AN
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Best Local Similarity
The invention relates to a murine pituitary-derived ligand polypeptide which is a ligand for the G-protein coupled orphan receptor designated GPR10 (human) or UHR-1 (rat). Cells transformed with a vector containing the ligand polypeptide encoding DNA are used to produce a recombinant ligand polypeptide. The ligand polypeptide, and its fragments, modulate function of the pituitary, central nervous system, pancreas and other tissues and can be used to screen for agents that modulate binding of the polypeptide to the receptor; to quantify the amount of receptor in a sample and to raise antibodies. They may also be used therapeutically, e.g. to treat senile dementia, Alaheimer's, Parkinson's or Huntington's diseases; Creutzfeld-Jakob disease; poisoning by heavy metals or drugs;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mutations or for isolation of related sequences, while the complete sequences may be used in gene therapy to provide wild-type protein. Antibodies specific for mutant or wild-type proteins are used as diagnostic reagents and for drug screening. The KCNQ2 and 3 proteins are useful in rational design of drugs and therapeutically (in replacement therapies). The forms of epilepsy associated with mutations in KCNQ2 and 3 sequences can now be diagnosed early (before symptoms are manifest), and better treatment options will be available. AAX57074-X57139 are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GPR10; UHR-1; modulator; pituitary; central nervous system; pancreas; tissue; screen; therapeutic; binding; senile dementia; ligand; murine; Alzheimer's disease; Parkinson's disease; Huntington's disease; drug; Creutzfeld-Jakob disease; poisoning; schizophrenia; growth hormone; secretion; diabetes; cancer; rheumatoid arthritis; epilepsy; vasopressor; gene therapy; transgenic animal; murine; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New polypeptide ligand for orphan G protein coupled receptors - used for treating disorders of central nervous system, pituitary and pancreas, an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hinuma
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                                                                                                                                                                                                                                                                                               Use of G protein-coupled receptor ligands - for modulating prosecretion or placental function, e.g. for treating menopausal tumours, autoimmune disease or abnormal pregnancy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         G protein-coupled receptor; GPCR; hypoovarianism; gonecyst cacogenesis; menopausal syndrome; euthyroid; hypometabolism; lactation; modulation; pituitary adenomatosis; brain tumour; emmeniopathy; autoimmune disease; prolactinoma; infertility; impotence; amenorrhea; galactorrhea; acromegaly; Chiari-Frommel syndrome; Argonz-del Castilo syndrome; Forbes-Albright syndrome; lymphoma; Sheehan syndrome; dyzzoospermia; contraceptive; placental function; choriocarcinoma; hydatid mole; abortion; unthrifty fetus; abnormal saccharometabolism; abnormal lipidmetabolism; oxytocia; prolactin secretion; PCR primer; ss
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Pred. No. 8
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PCR primers AAX15531-33 were used to amplify cDNA encoding a rat typ ligand polypeptide. The specification describes an agent for modulate prolactin secretion which comprises a ligand polypeptide or a salt, G protein-coupled receptor (GPCR) protein. The agents for promoting prolactin secretion can be used for treating or preventing enthyroic hypoporarianism, gonecyst cacogenesis, menopausal syndrome, enthyroic hypoproteins.

encoding a rat type n agent for modulating

syndrome,

gonecyst cacogenesis, menopausal syndrome, They can by used for promoting lactation i

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euthyroid

thyroid or domestic

Example

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Page 103; 241pp; English.

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RESULT 653
AAX36878/c
ID AAX368
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Best Local S
Matches 18
This sequence is a primer for the human XLIS gene of the invention. The XLIS fragments may be used to detect abnormalities in the expression of the XLIS transcripts or to compare their sequence with that of the XLIS transcripts from patients for in vitro especially prenatal diagnosis of lissencephaly (LIS) (or agyria-pachygyria), subcortical laminar neurodegenerative diseases such as Alzheimer's disease. These disorders mainly affect females as the XLIS gene is X-linked. The XLIS fragments may also be used to administer to patients to prevent or treat the above disorders and may be used as a tool in genetic counselling.

Oligonuclectides which bind to the fragments may be used to amplify the XLIS gene from a sample for comparison to normal samples in the in vitro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mammal and as an aphrodisiac. The agents for inhibiting prolactin secretion can be used for treating or preventing pituitary adenomatosis, brain tumour, emmeniopathy, autoimmune disease, prolactinoma, infertility, impotence, amenorrhea, galactorrhea, acromegaly, Chiari-Frommel syndrome, Argonz-del Castilo syndrome, Forbes-Albright syndrome, lymphoma, Sheehan syndrome or dyszoospermia. The inhibitory agents can also be used as contraceptives. The agents for modulating placental function can be used for treating or preventing choriocarcinoma, hydatid mole, irruption mole, abortion, unthrifty fetus, abnormal saccharometabolism, abnormal lipidmetabolism or oxytocia
                                                                                                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                                                    New gene and its gene product expressed in the brain, useful for diagnosing and treating disorders such as lissencephaly and subcortical laminar heterotopia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          XLIS gene, human; detection; diagnosis; prenatal diagnosis; therapy; lissencephaly; LIS; agyria-pachygyria; subcortical laminar heterotopia; SCIH; cortical dysgenesis; cryptogenic epilepsy; neurological discader; neurologenerative disease; Alzheimer's disease; X-linked disorder;
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Pred. No. 8
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                                                                                     PCR primers AAZ36052-Z36053 are used to amplify the mouse MLC2a gene. The primers and the PCR product are used in the method of the invention which relates to a novel quantitative reverse transcriptase polymerase chain reaction (RT-PCR) assay for quantitative gene expression. The method is used for determining a quantitative measure of the expression of a gene of interest in a biological sample by determining a normalised RNA prevalent for the gene of interest. The invention also relates to a method for determining the effect of a treatment on a quantitative measure of the expression of a gene of interest, or of a panel of gene of interest, in a sample by determining a normalised RNA equivalent for the gene of interest in a first untreated sample and a second treated sample. The methods are used for quantitative gene expression, where determination of changes in gene expression provides a measure of the biological response to a treatment or drug. The method have uses in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnosis regime. This may also be performed by amplifying XLIS cDNA frequency the mRNA in the sample. Antibodies to XLIS may be used to detect XLIS in a biological sample or can be administered to patients to prevent or treat the above disorders. They may also be used to putients to prevent or treat the above neurological disorders. In addition XLIS may be used to reat the above neurological disorders. In addition XLIS may be used as a marker of neuronal cells at an early stage of development; its discovery increases understanding of both the neuronal movement which leads to development of the cortical region of the brain and of the pathogenesis of the group of neuronal disorders mentioned above
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Quantitative analysis of gene expression using
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Pred. No. 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RT-PCR assays
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RESULT 656
AAF60520
ID AAF605
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AAC70378
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Best Local Similarity
Watches 18; Conserv
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                                                                                                                        Query Match
Best Local
                                                                                                              Matches
                                                                                                                                                                                           The present invention is concerned with a number of human single nucleotide polymorphisms (SNPs) which the inventors identified in human genes. These SNPs can be used in disease diagnosis and prediction of an individual's susceptibility to disease, in forensic and paternity testing and in genetic mapping. In particular, the SNPs of the invention can be used to diagnose susceptibility to diseases of the cardiovascular, endocrine and neurological systems, such as coronary artery disease, gchizophrenia, cancer, autoimmune diseases, Alzheimer's and Parkinson's
 AAF60520 standard;
                                                                                                                                                          Sequence 22 BP; 8 A; 3 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                 genetic analysis.
                                                                                                                                                                                                                                                                                                                                           Nucleic acid selected from one of 106 genes comprising single nucleotide polymorphisms, allele-specific oligonucleotides to the genes are useful for phenotypic correlations, forensics, paternity testing, medicine and
                                                                                                                                                                                                                                                                                                                                                                                                                    Altshuler D,
Lipshutz RJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-MAR-2000; 2000WO-US008440.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                neurological system; forensic testing; paternity testing; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single nucleotide polymorphism; SNP; human; genetic disease; disease susceptibility; cardiovascular system; endocrine system;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Single nucleotide polymorphism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         09-FEB-2001
                                                                                                                                                                                                                                                                                                          Claim 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        (WHED )
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                                                                                                             18;
                                                                                                                         Similarity
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                                                                                   TGTGGACCAGTATCGAGAGCAG 4554
                                                                                                                                                                                                                                                                                                         Fig 5; 214pp; English
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                                                            TTTGGACCAGTATAGACAGAAG 22
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nilarity 81.8%;
Conservative
                                                                                                             Conservative
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Patil N,
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 DNA;
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                                                                                                                      0.3%;
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klar P;
ВP
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Pred.
                                                                                                                        Score 15.6;
Pred. No. 8
                                                                                                             Mismatches
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No. 8.
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3.3e+02;
                                                                                                                       .3e+02
                                                                                                                                     DB 1;
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AAH45105
ID AAH4511
XX AAH451
XX AAH451
XX AAH451
XX AAH451
XX Diseas
XX Diseas
KW Down's
KW amyotn's
KW amyotn's
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                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a transgenic organism comprising a Plderived artificial chromosome (PAC) vector, which in turn comprises a pituitary adenylate cyclase-activating polypeptide (PACAP) receptor (PACR) gene. The present sequence is a PCR primer for human PACAP type 1 receptor (PAC 1R). PACAP is a neuropeptide which is widely expressed in the brain, and in various peripheral organs. PAC 1R is selective for PACAP and is expressed at high levels in the brain. PAC_IR is useful for preparing a medicament in the treatment and/or modulation of disturbances in stroke and other cerebrovascular diseases, cardiovascular diseases, leishmaniasis, immunosuppressive disorders, nociception (reaction to pain
                                                                                                                                                                                                                                                                                                                                                                    Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; PAC_1 receptor; transgenic organism; PACAP; brain; stroke; memory; pituitary adenylate cyclase-activating polypeptide receptor; nociception; PACAP type 1 receptor; PAC_1R; neuropeptide; cerebrovascular disease; cardiovascular disease; leīshmaniasis; immunosuppressive disorder;
Down's syndrome; cancer; neurodegenerative disease; Parkinson's amyotrophic lateral sclerosis; Huntington's disease; spinocerebellar ataxia-3; multiple sclerosis; diabetes mellitus
                                                                                                                                                                       AAH45105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene.
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                                              Disease detection; age-related disease; Alzheimer's disease;
                                                                          Beta-amyloid precursor protein,
                                                                                                            06-SEP-2001
                                                                                                                                         AAH45105;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 41; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New transgenic organism, useful for studying regulation of human P1-derived artificial chromosome (PAC) 1R gene expression, comprises a Prector or pituitary adenylate cyclase-activating polypeptide receptor
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                                                                                                                                                                                                                                                                                                                                                                                                    sensation) and learning and memory functions
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                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                   TCACCTCATTATGGAACTCTGG 4770
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                                                                                                                                                                                                                                                  TCACCTGCTTGTTGAACTCTGG 22
                                                                                                                                                                                                                                                                                                                                                                        BP; 3 A; 6 C; 5 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                          (first
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                                                                                                                                                                                                                                                                                                                           Score 15.6;
Pred. No. 8.
                                                                            APP,
                                                                                                                                                                                                                                                                                                              Mismatches
                                                                          coding sequence
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                                                                              PCR
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                                                                              primer.
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                                 disease;
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RESULT 658
AAF98936/c
ID AAF989
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AC AAF989
XY
T12-JUN
XX
Immuno
XX
Immuno
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Vaccin
KW Vaccin
KW Immuno
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a method for detecting a disease or pathological condition associated with molecular misreading of coding sequences in the genome and/or associated with the clearance of aber:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting a disease or pathological condition, associated with secretion of aberrant protein e.g. age-related diseases, by determining level of aberrant protein and/or its detectable part in body fluid and/or tissue.
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                                                                 Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma;
                                                                                                                                                                                                                                                                                                                                      AAF98936;
                                                                                                                                                                                                                                                                                                                                                                                                       AAF98936 standard; DNA;
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beta-amyloid precursor protein; APP; PCR primer; ss.
                                                                                                                                                                                                   Immunostimulatory nucleic acid #52.
                                                                                                                                                                                                                                                                     12-JUN-2001
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                                  allergy;
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                                     immune
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                                  deficiency; phosphorothicate;
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulatory nucleic acid. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a response and to activate immune cells. Note: the
                                                                                                     tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy, retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial anglogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25-SEP-199; 99US-0156113P
27-SEP-1999; 99US-0156135P
23-AUG-2000; 2000US-0227436P
                                                     Synthetic
                                                                                                                                                                                                             Anglogenesis inhibitor; ss; anglogenesis; solid tumour
                                                                                                                                                                                                                                                Angiogenesis inhibitory oligonucleotide #61.
                                                                                                                                                                                                                                                                                     13-DEC-2002
                                                                                                                                                                                                                                                                                                                        ABS77577
                                                                                                                                                                                                                                                                                                                                                         ABS77577 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 39; 338pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vaccinating against tumors, infectious diseases, allergies and asthma using immunostimulatory Py-rich and TG nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Krieg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  25-SEP-2000; 2000WO-US026383
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200122972-A2.
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                                                                                        scleroderma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (IOWA ) UNIV IOWA RES FOUND (COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5404 AAAAAGAAAAATGAAAATAAA 5425
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                                                                                                                                                                                                                                                                                                                                                                                                                                                22 AAAAACAAAAAAACAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Schetter C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 0 A; 0 C; 2 G; 20 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                     (first entry)
                                                                                      hypertrophic scar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
                                                                                                                                                                                                                                                                                                                                                           22
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                                                                                                                                                                                                                                                                                                                                                           ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.6;
Pred. No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                             growth;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22;
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WO200253141-A2

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RESULT 660
ABQ93268/c
ID ABQ932
XX ABQ932
XX ABQ932
XX 29-AUG
DT 29-AUG
DT 21-OCT
XX Micros
XW Micros
XW Polymo
KW Polymo
KW Giseas
KW Peneti
XX Aegilc
OS Tritic
XX Aegilc
OS Tritic
XX EP1217
XX EP1217
XX Z2-DE(
XX 22-DE(
XX 21-OE(
XX 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to inhibiting angiogenesis in a subject, comprising administering at least one antiangiogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glacom, retrolental fibroplasia, which is the property of the second contains of the property of the second corneal graft rejection, neovascular glacom, retrolental fibroplasia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Microsatellite marker; wheat; D genome; mapping; genotyping; polymorphism; phenotypic trait; QTL; quantitative trait locudisease-associated gene; development factor; quality factor; resistance factor; wheat product; identification; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound gramulation, intestinal adhesions, atherosclerosis, scleroderma hypertrophic scars. The present sequence is an antiangiogenic nucleic acid of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     genetically modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T. tauschii/wheat D genome microsatellite cfa2183 right PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-AUG-2003
21-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABQ93268 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 22 BP; 0 A; 0 C; 2 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                          22-DEC-2000; 2000EP-00403659
                                                                                                                                                                                                                                                                                                                                                                                 Triticum aestivum.
                                                                                                                                                                                                                                                                                                                                                                                                                     Aegilops tauschii.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABQ93268;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-566690/60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-DEC-2000; 2000US-025534P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-DEC-2001; 2001WO-US048458
                                          (INRG ) INRA INST NAT RECH AGRONOMIQUE
                                                                                                       22-DEC-2000; 2000EP-00403659
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (COLE-) COLEY PHARM GROUP INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5404 АЛАЛАGЛАЛАЛАТGAAЛATAAA 5425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAAAACAAAAAAACAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    81.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.6; DB 1
Pred. No. 8.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 °,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and
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RESULT ABV7717

ABV77172 standard;

DNA;

22

뫄

片 S

22

GCAGCTAATGAAGGAGAAGGAA 1

Query Match
Best Local Similarity
Matches 18; Conserv

2115 GCAGCAGATGAAGCGGAAGGAA 2136

Conservative

0; Pred. Score 15.6; Pred. No. 8. Mismatches

81.8%; 0.3%;

.3e+02

0;

Gaps

<u>,</u>

DB 1; Length 22; Indels

Sequence 22 BP; 2 A; 8 C; 2 G; 10 T; 0 U; 0 Other;

to standardise OS

field)

BXABXXXX

ABV77172;

27-OCT-2003 28-MAR-2003

(revised) (first entry)

Primer and probe for foot and mouth disease virus (FMDV)

a B

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The invention relates to a map of the bread wheat D genome comprising the CC genome location of a microsatellite marker selected from a group of 185 such markers (AB092713-AB092917). The invention also encompasses the use of left (AB092918-AB093102) and right (AB093103-AB093287) primers to aphilify and detect the microsatellite markers, and to identify genes responsible for a phenotypic trait of interest in wheat. Wheat is an allohexaploid species consisting of 3 diploid genomes designated A, B and CC allohexaploid species consisting of 3 diploid genomes designated A, B and CC different species. The D genome is thought to have been introduced in the CC tauschi (DD), probably involving only a limited number of genotypes of the protection of the protection of the genome, the large size of its genome, cand its low level of polymorphism, the genetic mapping of wheat has to the polymorphism makes them especially suitable for the genetic mapping of wheat has to comply the protection of the genetic mapping of wheat in a comply due to polymorphism makes them especially suitable for the genetic mapping of polymorphism makes them especially suitable for the genetic mapping of inheritance. The 185 microsatellite markers of the invention are developed from the ancestral diploid donor species Triticum tauschi and map to the wheat D genome, which is less polymorphise than the A or B map to the wheat not notably QTLs (quantitative trait loci). In CC particular they may be used for analysing genes and alleles implicated in CC disease and for identifying development factore, quality factors and feature, may be used for analysing genes and alleles implicated in mapping and genotyping diploid and polyploid species of Triticum comprises genetically modified wheat. The also be used in mapping and genotyping diploid and polyploid species of Triticum comprises genetically modified wheat. The present sequence represents a specifically claimed Triticum tauschi/wheat genome D concreases wheat or a related species; to assess whether or n
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Map of wheat D genome comprising the genome location of a microsatellite marker, useful for e.g. identifying genes responsible for a desired phenotypic trait, especially quantitative trait loci in wheat, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bernard M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 10; 105pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sourdille
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RESULT 662
ADA00879/c
ID ADA008
XX
AC ADA008
XX
AC ADA008
XX
MOUSE
XX
MOUSE
XX
ESDN;
KW ESDN;
KW ESDN;
KW ESDN;
KW TAMMUNE
KW IMMUNE
KW 
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Best Local Similarity 81.8%;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABV77170-83 represent primers and probes for foot and mouth disease virus (FMDV) 3D gene. A highly conserved area within the RNA polymerase gene is referred to as 3D. The probes and primers are useful for detecting a FMDV infection in a patient or for detecting an infection caused by any of several serotypes of FMDV. The primers and probes are especially useful for identifying FMDV infection caused by any of the FMDV serotypes such as Asia, A, C, O, Sat 1, Sat 2 or Sat 3. (Updated on 27-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel
virus
                                                                                                                                                                                                                                                                                                                                                                 ss; RT-PCR; type-I transmembrane protein; endothelial and smooth muscle cell-derived neuropilin-like molecule; ESDN; arterial sclerosis; restenosis;
        27-DEC-2001; 2001JP-00397725
                                                                                                                                                                        US2003129697-A1
                                                                                                                                                                                                                                                                                                          percutaneous transmembrane coronary angioplasty; PTCA; immune de immune disorder; tumour metastasis; diabetic retinopathy; mouse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse ESDN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADA00879 standard; DNA; 22 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Callahan JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20-MAY-2002; 2002WO-US015826.
                                                                                                                   10-JUL-2003
                                                                                                                                                                                                                               Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   primer; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FMDV; 3D gene; R
FMDV Asia; FMDV
                                                         10-JUL-2002; 2002US-00191436.
                                                                                                                                                                                                                                                                                         reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (TETR-) TETRACORE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-MAY-2001; 2001US-0291636P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200295074-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2003-129441/12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Page 8; 43pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  isolated nucleic acid useful for detecting foot and mouth disease infection in patient, and for detecting infection caused by ypes of the virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                μ.
                                                                                                                                                                                                                                                                                   transcriptase; primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTGGGTTTTATAAACCTGTGAT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CTGGGATTTATAAAACTGGGTT 403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RT-PCR primer #1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 5 A; 3 C; 5 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nelson
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A;
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Pred. No. 8.3e-
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 15.6; LL
No. 8.3e+02;
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                                                                                                                                                                                                                                                                                                                                          immune deficiency;
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RESULT 663
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a substantially purified form of a type-I transmembrane proteins, designated endothelial and smooth muscle cell-derived neuropilin-like molecule (ESDN). The ESDN polypeptide, cDNA, antibody or pharmaceutical composition is useful for treating arterial sclerosis, or restenosis after percutaneous transmembrane coronary angioplasty (PTCA). The ESDN polypeptide is also useful for screening reagents with antagonistic or agonistic activity. The antibody may be used for measuring the quantity of the ESDN polypeptide. The polypeptide is also useful for treating or preventing immune deficiencies and disorders, tumour metastases and diabetic retinopathy. The present sequence represents the mouse endothelial and smooth muscle cell-derived neuropilin-like molecule, ESDN, reverse transcriptase (RT)-PCR primer #1.
Treating non-allergic inflammatory diseases, such as psoriasis, allergic contact dermatitis, latex dermatitis or inflammatory be disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antilcer; gene therapy; vaccine; non-allergic inflammatory disease psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative collitis; Crohn's disease; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transmembrane coronary angioplasty.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Neuropilin-like molecule (BSDN) polypeptides or genes, useful for treating arterial sclerosis or restenosis after percutaneous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New human, mouse or rat Endothelial and Smooth muscle cell Derived
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                                                                                                                                                                                                                                              ) KRIEG A M.
) BERG D J.
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                                                                                                                                                                                     Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 2 A; 12 C; 1 G; 7 T; 0 U; 0 Other;
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Pred. No. 8.3e+02;
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RESULT 664
ADB36438/c
ID ADB364
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XX ADB364
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XX ADB364
AC ADB364
XX Immuno
XX ds; al
XM hypo-r
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Best Local Similarity
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Best Local S
                                         Matches
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                                                                                                                                                         The invention relates to a method of treating or preventing allergy or asthma which comprises administering to a subject a poly-G nucleic acid in an aerosol formulation. The method compositions of the present invention are useful for diagnosing and/or treating asthma and allergy especially in a hypo-responsive subject. The present sequence represent an immunostimulatory nucleic acid of the invention.
                                                                                                                                                                                                                                                                                                                                           Treating and/or preventing allergy or asthma using an immunostimulatory nucleic acid alone or in combination with an asthma/allergy medicament.
                                                                                                                     Sequence 22 BP; 0 A; 0 C; 2 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                       Claim 10; Page 6; 221pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Bratzler RL,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ds; allergy; asthma; poly-G nucleic acid; aerosol formulation;
hypo-responsive subject; immunostimulatory.
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(FOUR/)
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  5404 AAAAAGAAAAAATGAAAATAAA 5425
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FOURON Y.
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                                                          Score 15.6; DB 1;
Pred. No. 8.3e+02;
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Pred. No. 8.3e+02;
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                                       Mismatches
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                                                                             DB 1;
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                                                                                                                                                 RESULT 666
ADF44288
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Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                     Detector for identifying human papilloma virus subtypes, comprises carrier having two parts carrying first and second oligonuclectides that respectively hybridize with DNA contained in first and second subtypes of the virus.
                                                                                                                                                                                                                                                                              Sequence 22 BP; 2 A; 10 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                subtypes of human papilloma virus (HPV) contained in a sample. The oligonucleotides of the invention are useful for simultaneously detecting and identifying subtypes of HPVs. The present DNA sequence represents an HPV detection oligonucleotide of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probe; human papilloma virus; HPV; detection; identification; ss
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          JP2002360271-A
                           Human papillomavirus.
                                                                        HPV MM8
                                                                                                                  ADF44288;
                                                                                                                                                                                                                                                                                                                                                                 Claim 4; SEQ ID NO 616; 221pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2003-432398/41.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human papillomavirus.
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                                                detection; human papillomavirus;
                                                                                            12-FEB-2004
                                                                                                                                      ADF44288 standard; DNA; 22
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                                                                                                                                                                                                                                                                                                                                            The invention comprises oligonucleotides for detecting and identifying
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Shih C,
                                                                       detecting probe MM809.
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                                                                                            (first entry)
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Yeh C,
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                                                                                                                                                                                                                                   Score 15.6; DB 1;
Pred. No. 8.3e+02;
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                                                   HPV subtype; probe;
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RESULT 667
ADG76036/c
ID ADG760
XX ADG760
XX ADG760
XX Non-Cp
XX Non-Cp
XX B8; non-Cp
XX B8; no-Cp
XX B8; no-Cp
XX P1 asma
XX I1-DEC
XX 30-MAY
XX IMMU-
XX IMMU-
XX IMMU-
XX ILOPEZ
XX WPI; 2
XX PT New in
PT New in
PT New in
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XX INGELANCE
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                  ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes a novel detecting apparatus for identifying the subtypes of human papillomaviruses (HPV) contained in a sample which comprises a carrier which can load sample, a first oligonucleotide loaded on first part of the carrier and a second oligonucleotide loaded on second part of carrier, in which first and second oligonucleotides second part of carrier, in which first and second oligonucleotides hybridise with the DNA of the first and the second HPV subtype and can identify HPV subtype contained in sample at the same time. ADF43644-ADF44289 represent oligonucleotide probes used in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A detecting apparatus and a detecting method for identifying the subtypes of many species of human papilloma viruses at the same time and a composition for the detection.
              New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g
                                                                                                                                                                                                                               30-MAY-2003; 2003WO-EP005691.
                                                                                                                                                                                                                                                                                                  WO2003101375-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Non-CpG DNA oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADG76036;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADG76036 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 22 BP; 2 A; 10 C; 9 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 645; 166pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2003-600935/57.
                                                                                                                                                          (IMMU-) IMMUNOTECH SA
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                                                                                       2004-053333/05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGGGGCCGCCGCCAAGCCT 22
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                                                                                                                                                                                            2002CA-02388049
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Pred. No. 8.3e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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This invention
                                               Example 17; Page 81; 139pp; English
                                          relates to
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contain a non-palindromic sequence motif. Specifically, it refers to DNA oligonucleotides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune response is characterised by the proliferation, differentiation, cytokine and antibody production in B-cells, as well as cell differentiation of cytokine production in plasmacytoid dendritic cells. The present invention describes immunomodulator compositions that also comprise an antigen selected from, for example, viruses, bacteria, parasites, tumour cells and glycolipids. As such, these DNA oligos can be used in gene therapy for inducing B-cell activation, treating, preventing or ameliorating an immune system disorder or a tumoural disease including chronic myelogenous leukaemia, melanoma, Kaposi's sarcoma, and renal cell carcinoma. This oligonucleotide sequence is a non-cpG DNA oligo of the novel immunostimulatory oligonucleotides that

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Sequence 22 BP; 0 A; 0 C; 2 G; 20 T; 0 U; 0 Other;
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Query Match Best Local Similarity Conservative 81.8%; 0.3%; 0; Mismatches Pred. Score 15.6; Pred. No. 8 .3e+02 DB 1; Length 22; 0 0

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                      5404 AAAAAGAAAAAATGAAAATAAA 5425
22
AAAAACAAAAAAACAAAAAAAA 1
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RESULT 668
ADG76002/c
ID XX
ADG76002;
XX
AC ADG76002;
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DT 11-MAR-2004 (first
XX
DE Non-Cpg DNA oligonuc
XX
DE PTON-Cpg DNA oligonuc
XX
DE NON-Cpg DNA Oligonuc
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DNA OLIGONUC
XX
DE SAMPLE 17; Page 80;
XX
DR Example 17; Page 80;
XX
This invention relation
CC contain a non-palinc
CC response in animals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ss; non-CpG; immunostimulatory; non-palindromic; immune response; proliferation; differentiation; cytokine; antibody production; B-plasmacytoid dendritic cell; immunomodulator; gene therapy; chronic myelogenous leukaemia; melanoma; Kaposi's sarcoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Non-CpG DNA oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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30-MAY-2003; 2003WO-EP005691.

(IMMU-) IMMUNOTECH SA

30-MAY-2002; 2002CA-02388049.

New immunostimulatory oligonucleotide comprising non-palindromic nucleic acid sequence motif, useful for inducing B-cell activation, treating, preventing or ameliorating immune system disorder or tumoral disease e.g.

139pp, English

This invention relates to novel immunostimulatory oligonucleotides that contain a non-palindromic sequence motif. Specifically, it refers to DN oligonucleotides (without a CpG motif), which can stimulate an immune response in animals of the order of primate, including humans. The immune DNA

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ARESULT 669
ADH70416/c
ID ADH704
AC ADH706
AC 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; degenerative nervous system disease; multiple sclerosis; lzheimer's disease; hypersensitivity; depolasture's syndrome; allergy; type II hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity, leprosy; infectious disease; viral infective; fungal infection; Candida; parasitic infection; schistosome; filaria; bacterial infection, Mycobacterium; neoplastic disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease, hypersensitivity disease; infectious disease; neoplastic dis
                                                                                                                                                                                                                                                             Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, comparation of the control o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22
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19-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          тосат
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              2004-059052/06.
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95US-00531241.
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Pred. No. 8.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 T; 0 U;
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associated diseases

which

a kit for diagnosing a h comprises a panel of comprises a

panel

and treating f nucleic acid

acid

The present invention describes a compound 8-80 nucleobases in targeted to, and which specifically hybridises with a nucleic a molecule encoding perilipin, and inhibits the expression of per Also described: (1) a compound 8-80 nucleobases in length that specifically hybridises with at least an 8-nucleobase portion o active site on a nucleic acid molecule encoding perilipin; (2)

portion of an ipin; (2) a

perilipin

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The invention relates to

Disclosure;

SEQ

ID NO 610; 164pp; English

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RESULT 670
ADJ78305/c
ID ADJ783
XX ADJ783
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                      New compounds, particularly antisense oligonucleotides targeted to nucleic acids encoding perilipin, useful for treating a metabolic disorder e.g. obesity, diabetes or atherosclerosis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           perilipin; perilipin inhibitor; antisense oligonucleotide; antidiabetic;
anorectic; antiarteriosclerotic; cardiant; metabolic disorder; diabetes;
obesity; atherosclerosis; mouse; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse perilipin reverse PCR primer SEQ ID NO:13.
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Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
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    This invention relates to a novel method for the diagnosis of greater susceptibility to gastric carcinoma, comprising searching for a possible polymorphism in the promoter of the epithelial caderine (B-caderine) gene. The method is useful for identifying a genetic polymorphism that leads to a greater predisposition to the onset of gastric carcinoma. The method is relatively simple, quick, accurate and reliable. The present
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                                                                                                                                                                                                                                                                             Claim 11; Page 12; 17pp; English.
                                                                                                                                                                                                                                                                                                                                                 Identifying greater susceptibility to gastric carcinoma polymorphisms in the promoter of the E-caderine gene.
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llarity 81.8%;
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Pred. No. 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CMV; inverted polarity region;
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RESULT 673
AAQ30313
ID AAQ303
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This oligomer contains an inverted polarity region formed from an oxyloso dimer synthon. Residues 13 and 14 are linked via an oxyloso group (i.e. nucleotides that have xylose sugar linked via the oxylene ring). The sequence is designed to target the Cytomegalovirus beginning at nucleotide 176 and to covalently cross-link to it. See also AAQ20162-
                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligomer CMV802
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07-DEC-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New sequence-specific non-photo-activated crosslinking the major groove of duplex DNA and are esp. useful for infections e.g. HIV.
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                         modified_base
                                                                                                                                                                                                                                                                                                                                                                                                     Cytomegalovirus; herpes; hepetitis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ30313 standard;
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(first entry)
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91US-00640654.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     for forming triplex with CMV target duplex.
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/note= "N-methyl.
                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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                                                             *tag= c
mod_base= OTHER
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/mod_base= OTHER
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Dase= OTHER
המיחים N6 methyl-8-oxo
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                                                                                                                                    "OTHER=
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      Q.
                                                                                                                                                                                                                                                                                                                                                                                                     AIDS; modified; HIV; RSV; HPV; malignancy; ss.
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                                                N6 methyl-8-oxo 2' deoxyadenine"
                                                                                                                                  N6 methyl-8-oxo
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17-APR-1991;
17-APR-1991;
17-APR-1991;
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                    Froehler B,
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18-JAN-1991;
                                                                                                                                                   25-NOV-1991;
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                                         (GILE-)
1992-217083/26.
                                         GILEAD
                    Krawczyk
                                                                       91US-00686544.
91US-00686546.
91US-00686547.
                                         SCI INC.
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/note= "OTHER= N6 methyl-8-oxo 2' deoxyadenine"
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"see comments"
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                    Matteucci MD,
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                    Milligan
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New oligomers contg. modified bases - which form a triplex with G-C doublet in a DNA duplex, for treating and diagnosing HIV, hepatitis, herpes malignancy and inflammation.

Claim 12; Page 68; 77pp; English.

The synthetic oligomer is capable of forming a triplex at physiological pH with a purine rich target sequence by coupling into the major groove of the duplex. The specific target sequence of this oligomer is a cytomegalovirus duplex beginning at nucleotide 176 contg. a purine-rich region concentrated on one chain of the duplex. The oligomer, and others like it are useful in diagnosis and therapy of diseases characterised by specific DNA duplex targets, e.g. cytomegalovirus; HFV; HTV, hepatitis B, herpes, malignant tumours and inflammation. The triple helices form under mild conditions thus assays may be carried out without subjecting the test specimen to harsh conditions. The oligomer contains an inverted polarity region formed from an o-xyloso dimer synthon. The linking gp. is o-xyloso (nucleotides have the 3' positions of xylose sugars linked via the o-xylene ring). Two nucleotides are coupled through a xylene residue to form the dimer synthon. This additional modifications may render the

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RESULT 674
ANT12445/c
ID ANT124
XX ANT124
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Best Local S
Matches 16
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                                                                                                                                                                                                                                              AAT12435-T12454 represent phosphorothioate oligonuclectides with antiviral activity. These sequences, and the phosphorothioate oligonuclectides represented by AAT12418-T12434 (which are complementary to regions of the mRNA 4 or 5 of herpes simplex virus 1 (HSV)), are effective in the prevention and treatment of viral infection. The sequences are especially effective against infection by HSV, HIV or varicella zoster virus (VZV)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (KAKE )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JUL-1994;
01-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oligomer stable to nuclease activity. The oligomer is able to inhibit gene expression, as verified by in vitro systems. See also AAQ25452-25501 and AAQ30226-448. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Antiviral phosphoro:thioate oligo:nucleotide(s) - active against herpes simplex virus 1, HIV and varicella zoster virus.
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17
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                                                                                                Conservative
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94JP-00268603
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                                                                                                                     Pred. No. 8.6e+02;
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Pred. No. 8.6e+02;
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Query Match

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Length 17;

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ID VAV9
XX AAV9
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XX
                                             CC A method has been developed for the identification of a nucleic acid cc capable of modulating a process in a biological system. The method CC comprises: (a) introducing into the system a random library of nucleic CC acid catalysts (NAC) having a substrate binding domain (SBD), comprising CC a random sequence, and a catalytic domain (CD); and (b) identifying NAC CC in systems where modulation has occurred and/or determining the sequence CC of at least part of the SBDs in such systems. Nucleic acid molecules with CC endonuclease activity and catalytic activity, from the present invention, CC are used to modulate gene expression in plant and mammalian cells and to CC cleave target nucleic acid, particularly for treating systemic diseases CC cused by specific RNA, e.g. cancer, inflammation, psoriasis, non-hepatic CC with RNA-cleaving activity that modulate expression of the RAE gene, are used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or CC generally any condition associated with the level of c-raf. Introduction CC used to treat cancer, restenosis, psoriasis or rheumatoid arthritis, or CC generally any condition associated with the level of c-raf. Introduction of sugar/phosphate modifications increases stability against nuclease and CC activity. AAV90922 to AAV93877 represent NACs that can be used in the CC method, specifically for modulating the expression of a Raf gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           02-OCT-1997;
02-OCT-1997;
05-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Larget; substrate; catalyst; modulation; expression; Raf gene; delivery screening; identification; synthesis; deprotection; purification; cance: inflammation; psoriasis; non-hepatic ascites; infection; genetic drift; restenosis; rheumatoid arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying new catalytic nucleic acid that modulates selected processes - especially ribozymes that cleave Raf RNA for treating cancer, restenosis, and also new ribozymes and modified nucleoside triphosphates used as antiviral agents and synthons.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 177; Page 157; 259pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-JUL-1997;
22-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           675
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Beigelman L, I
J, Workman CT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matulic-Adamic J, Reyn
Beigelman L, Mcswiggen
, Workman CT, Beaudry
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97US-0064866P.
97US-0068212P.
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97US-0056808P.
97US-0061321P.
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97US-0049002P.
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ggen JA, Karpeisky A,
udry A, Sweedler D;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              delivery;
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RESULT 676
AAA25454/c
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                                                                                                           CC with a target sequence and contain at least one phosphoro (di) thicate (CC link, having endonuclease activity. (A), and more generally any catalytic mucleic acid (A') that modulates expression of the oestrogen receptor cc gene, are used to treat cancer (particularly of breast or endometrium), in vivo or by transforming cells ex vivo and implanting treated cells, or cc for other conditions associated with levels of oestrogen receptor. CC Because of the high selectivity for targeted RNA, (A) can also be used to correlate inhibition of gene expression with alterations in phenotype, cc particularly for identification of therapeutic targets, and as research cc used with DNA). The combination of modifications in (A) improves are used with DNA). The combination of modifications in (A) improves cc vasistance to nucleases, binding affinity and/or activity. AAA23503 to CAAA24747 represent oestrogen receptor hammerhead ribozyme sequences, and CC AAA24748 to AAA25992 represent their corresponding target sequences, and CC AAA2593 to AAA26107 to AAA26271 represent their corresponding target sequences and CC aAA24748 to AAA25105 to AAA26271 represent obstrogen receptor hairpin ribozyme sequences and AAA26219 to AAA26271 represent other ribozyme sequences and cr invention
Query Match
Best Local Similarity
Matches 16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thompson JD, Beig
Reynolds M, Zwick
Matulic-Adamic J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oestrogen receptor; c-raf; k-ras; bcl-2; ribozyme; cleavage; hammerhead ribozyme; hairpin ribozyme; antisense oligonucleotide; gene expression modification; cancer; phosphorothioate; endonuclease;
                                                                Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes nucleic acids (A) that interact
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 77; Page 79; 148pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New nucleic acids that interact, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-APR-1998;
23-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oestrogen receptor hammerhead ribozyme target sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          to treat cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2000-013248/01.
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                                                                  BP;
 0.3%;
ilarity 94.1%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zwick M,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Beigelman L,
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98US-00103636.
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                                                                   0 C; 1
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Score 15.4; DB 1; Length 17; Pred. No. 8.6e+02; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mcswiggen JA,
is T, Woolf T,
                                                                  G; 15 T; 0 U; 0 Other;
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1; Mismatches 1
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Haeberli P;
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RESULT 677
ABL46875/c
ID ABL468
XX ABL468
XX ABL468
XX Human;
XX Human;
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XX 12-FEE
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ABT38160
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XX ABT3
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                                                                                                                                                                                                             RESULT 678
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to oligonucleotides that downregulate the expression of human Grb2-related with Insert Domain (GRID) gene. GRID is a T-cell co-stimulatory adaptor protein. The oligonucleotides are useful for modulating the expression of GRID, to treat conditions such as tissue/graft rejection and leukaemia. The oligonucleotides can also be administered in conjunction with other therapies such as radiation, chemotherapy and cyclosporin treatment. The present oligonucleotide was used to illustrate the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 17 BP; 3 A; 5 C; 5 G; 0 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid(s) for regulating the Grb2-related with Ins (GRID) gene comprises using antisense and enzymatic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABL46875 standard; RNA; 17
                        Tumour suppression related human fukutin oligo SEQ ID No 3797.
                                                                        12-JUN-2003
                                                                                                                      ABT38160
                                                                                                                                                                  ABT38160 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Page 69; 108pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jarvis T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; Grb2-related with Insert Domain; GRID; T-cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human GRID G-cleaver ribozyme substrate oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              molecules such as hammerhead
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                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                                                                                                                                                                                                                                    GAGGCCTTCGTGAAACC
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                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                   94.1%;
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                                                                                                                                                                    ВP
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                                                                                                                                                                                                                                                                                                                                                                                0;
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                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4; DB 1; Length 17; Pred. No. 8.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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ABZ22872/
ID ABZ2
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                                                                                                                                                                                                                                                                                                                                                           밁
                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel isolated 17 mer nucleic acid sequence, cc given in the specification, a sequence containing at least 15 consecutive cnucleotides from the 17 mer sequence with, after optimal cc alignment, at least 80 % identity to the 17 mer sequence with, after optimal cc hybridizes to them under highly stringent conditions, or the complement cc fany of them, or the corresponding RNA. The novel isolated nucleic acids of the invention are useful as probes and primers for detecting, cc identifying, quantifying and/or amplifying a nucleic acid, e.g. as one cc component of a gene chip, in vitro as (anti)sense reagents, and for production of recombinant polypeptides. Any of the nucleic acids, colls containing the vector or antibodies directed against the polypeptides are useful for preparation of pharmaceuticals for prevention and/or treatment of viral cd degeneration, specifically cancer but also Alzheimer's disease and containing the schizophrenia. Analysis of the expression of the 17 mer nucleic acids in patient samples is useful for diagnosis and/or prognosis of these cc diseases. The polypeptides can also be used to generate antibodies, and components of these caids such the polypeptide and antibodies are useful as components of these characterises and contained the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene characterises and contained the polypeptide and antibodies are useful as components of protein chips. The nucleic acid sequences of the invention can be used in gene chips. The nucleic acid sequences of the invention can be used in gene characterises and contained the invention can be used in gene characterises and contained the invention can be used in gene characterises and contained the invention can be used in gene chips.
                                                                                                                                                                               RESULT 679
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
Phosphorothioate; locked nucleic acid; LNA; immunostimulatory; cytostatic; antimicrobial; gene therapy; pathogenic infection; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 477; 720pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cytostatic; virucide; neuroprotective; nootropic; neuroleptic; gene chip; antisense; sense; tumour; cell degeneration; cancer; Alzheimer's disease; schizophrenia; protein chip; gene therapy; tumour suppression; human fukutin; ds.
                                                                                  07-APR-2003
                                                                                                                  ABZ22872
                                                                                                                                                                                                                                                                                                                                                         Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-313353/30.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-SEP-2001; 2001FR-00011978
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                               Locked nucleic acid oligonucleotide LNA5.
                                                                                                                                               ABZ22872 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MOLE-) MOLECULAR ENGINES LAB
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                                                                                                                                                                                                                                                           3936 GATCAACCCACAGCACA 3952
                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                              GATCAACCCACAGAACA 17
                                                                                                                                                                                                                                                                                                                                                         BP; 8 A;
                                                                                                                                                                                                                                                                                            Conservative
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                                                                                (first
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                                                                                entry)
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                                                                                                                                                                                                                                                                                           Score 15.4; DB 1; Length 17; Pred. No. 8.6e+02; O; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                         0 Other;
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AC ADB4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cc comprising an oligonucleotide having at least one locked nucleic acid (C based on a functional molecy. Also described: (1) a complex (II) and a DNA sequence having a complementary sequence to the comprising (I) and a DNA sequence to the coligonucleotide, and encoding a gene under the control of a promoter; (2) a pharmaceutical composition (III) comprising (II) and a carrier or (C diluent; (3) a device loaded with (III); and (4) an oligonucleotide (IV) comprising a first region comprising an oligonucleotide sequence having at least one LNA, and a second region comprising an immunostimulatory (2) cligonucleotide region containing at least one unmethylated (C di-can be used in gene therapy. (I) and (II) can be used in medicament for the treatment or the prevention of (C the manufacture of a medicament for the treatment or the prevention of (C pathogenic infections or cancer. (I) is useful for the preparation of (C intention) and natigen or therapeutic protein, and formulating the resulting complex with a pharmaceutical carrier. The present sequence represents a complex with a pharmaceutical carrier. The present sequence represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel locked nucleic acid conjugate useful in manufacturing a medicament for treating or preventing pathogenic infections or cancer, has an oligonucleotide having locked nucleic acid based on a functional moiety.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Key
                                                                      Tumour suppression/reversion associated nucleotide #5701
                                                                                                                                                                               ADB45378;
                                                                                                                                                                                                                           ADB45378 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a locked nucleic acid (LNA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 1; Page 20; 101pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            invention
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16; Conserv
                                                                                                                                                                                                                                                                                                                                                            GAGAGAGAGAGAGAG
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/mod_base= OTHER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.4; DB 1;
Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 17;
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cytostatic; antiviral; neuroprotective; nootropic; neuroleptic; primer; probe; tumour suppression; tumour reversion; apoptosis;

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ADIS1815
ID ADIS18
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XX TIS-APR
XX TUMOUT
KW CYCOST
KW CYCOST
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Analysis of the expression of the nucleotides can be used for diagnosis and/or prognesis of these diseases. The nucleotides and polypeptides can also be used to screen for their specific interactive molecules.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                            cytostatic; virucide; neuroprotective; nootropic; neuroleptic; probe; primer; PCR; gene chip; antisense; viral disease; tumour;
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Homo sapiens
                                                                                                                                                         tumour suppression; tumour reversion; apoptosis; virus resistance;
                                                                                                                                                                                                                                 Human
                                                                                                                                                                                                                                                                                                  15-APR-2004
                                                                                                                                                                                                                                                                                                                                                                   ADI51815
                                                                                                                                                                                                                                                                                                                                                                                                                                    ADI51815 standard; DNA; 17 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               potentially useful for treating diseases associated with abnormal expression of the nucleotides.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17-SEP-2001; 2001FR-00011981.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-MAY-2003
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                                                                                                                                                                                                                                 tumour
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                                                                                                                                                                                                                              suppression/reversion-related DNA sequence SeqID4318
                                                                                                                                                                                                                                                                                               (first entry)
                                                                cancer; Alzheimer's disease; schizophrenia; ds; human.
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in the phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses. The invention may be useful for the development of compounds with a cytostatic, virucide, neuroprotective, nootropic or neuroleptic activity. The DNA sequences may be useful as probes and primers for detecting, indentifying, quantifying and/or amplifying nucleic acid, for example as one component of a gene chip, in vitro as antisense reagents and for production of recombinant polypeptides. The invention may therefore be useful for preparation of pharmaceuticals for prevention and/or treatment of viral diseases that are characterised by development of tumours or cell degeneration, specifically cancer but also Alzheimer's disease and schizophrenia. The present sequence is that of a nucleic acid sequence of the invention. Note: The sequence data for this patent did not form part of the printed manification by the printed sequence of the printed manification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      New isolated nucleic acid, useful for treating viral diseases associated with tumors and cell degeneration, also related polypeptides, antibodies and transfected cells.
Sequence 17 BP; 8 A; 6 C; 2 G; 1 T; 0 U; 0 Other;
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                                        at ftp.wipo.int/pub/publishedpct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; SEQ ID NO 4318; 30pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-313354/30.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Telerman A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MOLE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-SEP-2002; 2002WO-IB004523.
                                                               specification,
                                                                                                                                                                                                                                                                                                                                                                       invention relates to novel isolated nucleic acid sequences involved
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULAR ENGINES LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Amson R,
                                                               but was obtained in electronic format
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tuijnder M;
                                                               part of the printed directly from WIPO
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ADM54233/c
ID ADM542
XX ADM542
XX ADM542
XX ADM542
XX Human
XX Hum
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Best Local S
Matches 16
                                             10-FEB-2000; 2000US-0181594P
                                                                                                                               23-FEB-2001; 2001US-00792818.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; ss; GRID; Grb2-related with insert domain; hammerhead ribozyme; NCH ribozyme; G-cleaver ribozyme; Zinzyme; DNAzyme; amberzyme; Inozyme hairpin ribozyme; tissue rejection; graft rejection; leukaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADM54233;
                                                                                                                                                                                                                          17-JUL-2003
                                                                                                                                                                                                                                                                                                            US2003134806-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human GRID mRNA substrate sequence #508.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADM54233 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3936 GATCAACCCACAGCACA 3952
                                                                                                                                                                                                                                                                                                                                                                                                   sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GATCAACCCACAGAACA 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15.4; DB 1; Length 17; Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Inozyme;
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RESULT 683
ADH70294/c
ID ADH702
XX ADH702
XX ADH702
XX ADH702
XX Human
XX Human
XX Hypers
KW Hypers
KW Aller
KW Aller
KW Aller
KW HILL
KW Hypers
KW HILL
KW Hypers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to a nucleic acid molecule that down-regulates
CC expression of Grb2-related with insert domain (GRID) gene, e.g. a
CC hammerhead ribozyme, NCH ribozyme, G-cleaver ribozyme, Zinzyme, DNAzyme,
CC hammerhead ribozyme or hairpin ribozyme. Also include are a mammalian cell
CI including the novel nucleic acid molecule, reducing GRID activity in a
CC cell by contacting the cell with the novel nucleic acid molecule,
CC treating a patient having a condition associated with the level of GRID
CC (e.g. tissue/graft rejection or leukaemia) by contacting the cell with
CC the novel nucleic acid molecule, cleaving RNA of a GRID gene by
CC contacting the cell with the novel nucleic acid molecule, an expression
CC vector comprising a nucleic acid sequences (encoding at least the novel
CC nucleic acid molecule in a manner that allows its expression), a
CC mammalian cell including the expression vector and an enzymatic nucleic
CC acid molecule that cleaves RNA derived from a GRID gene. The nucleic acid
CC molecule is useful for treating a condition associated with the level of
CC GRID, e.g. tissue/graft rejection and leukaemia. The present sequence is
CC a target region for the enzymatic nucleic acids of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                              human; T-cell associated disease; Vbeta; autoimmune disease; degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; hypersensitivity disease; neoplastic disease; addison's disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; alzheimer's disease; hypersensitivity disease; type I hypersensitivity; disease; type I hypersensitivity; Goodpasture's syndrome; allergy; type II hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; leprosy; infection; disease; viral infection; HIV; fungal infection; Candida; parasitic infection; schistosome; filaria; bacterial infection, Mycobacterium; neoplastic disease; lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New nucleic acid molecule with insert domain (GRID) associated with the level
                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADH70294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADH70294 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; SEQ ID NO 508; 74pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CARL/)
(MCSW/)
(HAMB/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               leukemia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JARV/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4429 GAGGCCTTGGTGAAACC 4445
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vbeta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17
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) CARLOWITZ I V.
) MCSWIGGEN J.
) HAMBLIN P A.
) ELLIS J H.
                                                                                                                               cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Carlowitz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       repeat sequence #84.
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C;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              that down-regulates expression of Grb2-related gene, useful for treating a condition of GRID, e.g. tissue/graft rejection and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mcswiggen J, Hamblin PA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 T; 4 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
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The invention relates to a kit for diagnosing and treating T-cell CC associated diseases which comprises a panel of nucleic acid primers CC specifically priming and allowing amplification of each Vbeta gene, CC vbetaRNA or cDNA. The kit is useful for diagnosing organ transplant CC rejection and diagnosing and treating T-cell associated diseases CC including autoimmune diseases, degenerative nervous system diseases, CC graft versus host diseases, hypersensitivity diseases, infectious diseases, CC and neoplastic diseases. Autoimmune diseases include Addison's disease, CC atrophic gastritis. Degenerative nervous system diseases include multiple CC atrophic gastrities buch as contact with allergens that lead to CC clarosis and Alzheimer's disease. Hypersensitivity diseases include Type CC in hypersensitivities such as those present in CC Goodpasture's syndrome and Type IV hypersensitivities such as those caused by viruses such as those caused by viruses such as those caused by classification and bacterial infections such as those caused by CC such as leukamias filaria and bacterial infections such as those caused by CC such as leukamias filaria and bacterial infections such as those caused by the prophastic diseases include lymphoproliferative diseases of the hair.
                                          such as lew
breast. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, nucleic acid primers specifically priming and allowing amplific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-SEP-1994;
19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO 488; 164pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2004-059052/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Vbeta gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (HOOD/) HOOD L E. (ROWE/) ROWEN L.
                                                                leukaemias,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Rowen L;
                                          present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94US-00309335.
95US-00531241.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-00263959
                                      lymphomas and cancers such as cancer of the brain, sequence represents a Vbeta gene repeat sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprises
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B

Sequence 17 BP; 0 Ą, 8 C; 0 G; 9 T; 0 U; 0 Other;

밁 S Matches Best Query Match Local 1180 AGAGAAAGAGAGAGAGA 1196 17 Similarity Conservative 0.3%; 0 Score 15.4; DB 1; Pred. No. 8.6e+02; Mismatches DB 1; Length 17; 0 Gaps

0

ADH70390 standard; DNA; 17 ADH70390 ВP

Human Vbeta gene repeat sequence #180.

25-MAR-2004

(first entry)

RESULT 684
ADH70390/c
ID ADH703
XX ADH703
AC ADH703
XX Z5-MAR
AC LUMMAN
AC HUMMAN
KW HUMMAN
KW HUMMAN
KW HUMMAN
KW HYPERS
KW ADdiso
KW ADdiso
KW ALZhei
KW Alzhei
KW Allerg
KW HIY; f
KW Filari degenerative nervous system disease; graft versus host disease; hypersensitivity disease; infectious disease; neoplastic disease; hypersensitivity disease; atrophic gastritis; degenerative nervous system disease; multiple sclerosis; allzhelmer's disease; hypersensitivity disease; type I hypersensitivity; allergy; type II hypersensitivity; Goodpasture's syndrome; type IV hypersensitivity; leprosy; infectious disease; viral infection; fingal infection; Candida; parasitic infection; schiatosome; filaria; bacterial infection; Mycobacterium; neoplastic disease; human; T-cell associated disease; Vbeta; autoimmune dis

US2002150891-A1

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RESULT 685
ADH70382/c
ID ADH703
XX ADH703
AC ADH703
XX
DT 25-MAR
XX
DE Human
XX
DH Human;
KW degene
KW hypers
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                                                                                                                                                                                                                                                                                                                                                                         S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a kit for diagnosing and treating T-cell associated diseases which comprises a panel of nucleic acid primers of specifically priming and allowing amplification of each Vbeta gene, CC VbetaRNA or CDNA. The kit is useful for diagnosing organ transplant crejection and diagnosing and treating T-cell associated diseases. CC including autoimmune diseases, degenerative nervous system diseases, careful to result the control of the cont
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 16
degenerative nervous system disease; graft versus host di hypersensitivity disease; infectious disease; neoplastic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   lymphoproliferative disease; leukaemia; lymphoma; cancer; brain cancer; breast cancer; ds.
                                             human; T-cell associated disease; Vbeta; autoimmune disease;
                                                                                    Human Vbeta gene repeat sequence
                                                                                                                                   25-MAR-2004
                                                                                                                                                                               ADH70382
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acid primers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-SEP-1994;
19-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-MAR-1999;
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                                                                                                                                                                                                                       ADH70382 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Kit for diagnozing and treating T-cell associated diseases e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (HOOD/) HOOD L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (ROWE/) ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                           1181 GAGAAAGAGAGAGAGAG 1197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2004-059052/06
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                                                                                                                                                                                                                                                                                                                                     17
                                                                                                                                                                                                                                                                                                                                                                                                                      16;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rowen L;
                                                                                                                                                                                                                                                                                                                                     GAGAGAGAGAGAGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               degenerative nervous system and infectious disease, comprises primers specifically priming and allowing amplification of a
                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                 (first entry)
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95US-00531241.
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                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                           94.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        9 C; 0 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               584; 164pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.3%;
                                                                                                                                                                                                                                                                                                                                  -
                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15.4;
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                        #172.
                                                                                                                                                                                                                                                                                                                                                                                                                                           8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                      disease;
                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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RESULT 686
AAT32141
ID AAT321
XX
AC AAT321
XX

AAT32141 standard; DNA; 18

AAT32141;

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The invention relates to a kit for diagnosing and treating T-cell cc associated diseases which comprises a panel of nucleic acid primers cspecifically priming and allowing amplification of each Vbeta gene, the versus of the kit is useful for diagnosing organ transplant cc rejection and diagnosing and treating T-cell associated diseases, cincluding autoimmune diseases, degenerative nervous system diseases, cc graft versus host diseases, hypersensitivity diseases, infectious diseases, cc and neoplastic diseases. Autoimmune diseases include Addison's diseases, cc arrophic gastritis. Degenerative nervous system diseases include multiple cc atrophic gastritis such as contact with allergens that lead to cc allergies, Type II hypersensitivities such as those present in cc Goodpasture's syndrome and Type IV hypersensitivities such as those caused by viruses such as HIV, fungal infections such as those caused by caused by viruses such as HIV, fungal infections such as those caused by chistosomes, filaria and bacterial infections such as those caused by chistosomes, filaria and bacterial infections such as those caused by chistosomes, filaria and bacterial infections such as those caused by chistosomes, filaria and bacterial infections such as those caused by chistosomes, filaria and bacterial infections such as those caused by chistosomes, filaria and bacterial infections such as those caused by chistosomes, filaria and bacterial and bacterial and bacterial such as those caused by chistosomes, filaria and bacterial and bacter
                                                                                      Query Match
Best Local S
Matches 16
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                                                                                                                                                                                                                                                                             Sequence 17 BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; SEQ ID NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kit for diagnozing and treating T-cell associated diseases e.g. autoimmune, degenerative nervous system and infectious disease, nucleic acid primers specifically priming and allowing amplific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2004-059052/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hood LE, Rowen L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US2002150891-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Vbeta gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              breast cancer; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HOOD/) HOOD L E
(ROWE/) ROWEN L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-SEP-1994;
19-SEP-1995;
                                                                                                                                      Local Similarity
1180 AGAGAAAGAGAGAGA 1196
                                                                                      16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HOOD L E.
                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94US-00309335.
95US-00531241.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9908-00263959.
                                                                                                                                      0.3%;
                                                                                                                                                                                                                                                                             8 C; 0 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        576; 164pp; English.
                                                                                           <u>.</u>
                                                                                                                                           Score 15.4; DB 1;
Pred. No. 8.6e+02;
                                                                                                Mismatches
                                                                                                                                                                                  Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          disease, comprises amplification of a
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                                                                                           Gaps
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RESULT 687
AAT98144/c
ID AAT981
XX
AC AAT981
XX
DT 13-MAR
XX
DE Primer
XX
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                                                                                                                                                                                                                                                                                                                                     The present sequence is an example of a complementary sense strand from a novel DNA sequencing oligonucleotide called a "prinker" (primer/linker), which comprises a priming site, labelling region, cohesive end and complementary strand. The priming site is the optimal target for annealing prior to treatment with polymerase. The labelling region is a template sequence which directs DNA polymerase to incorporate multiple labelled, e.g. radioactive nucleotides. The cohesive end provides compatible ends for ligation of prinkers to restriction fragments. The complementary strand provides a region of double stranded DNA which is required by DNA ligases for the attachment of the prinker to a restriction fragment. A prefd. sequencing procedure comprises the generation of prinkers to the fragments sepn. and purificn. of prinker ligation of prinkers to the fragments, sepn. and purificn. of prinker to settached restriction fragments, conc. and buffer exchange, generation and sepn. of sequencing prods., exposure of X-ray film to sequencing prods.
                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New oligo:nucleotide(s) for DNA sequencing - having a priming site, labelling region and a cohesive end complementary to a restriction fragment sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-FEB-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sense strand; priming site;
                 Primer V-alpha(16) for T-cell receptor alpha chain
                                                                                                                                                                                                                                                                                                            Sequence 18 BP; 14 A; 2 C; 1 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 5; 23pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1996-105934/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   misc_feature
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                                                13-MAR-1998
                                                                              AAT98144;
                                                                                                           AAT98144 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Leonard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (AMIC-) AMICON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9602673-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA sequencing "prinker" (primer/linker) complementary sense strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-SEP-1996
                                                                                                                                                                                                            5401 ACAAAAAAGAAAAAATG 5417
                                                                                                                                                                                                                                                 al Similarity
16; Conserv
                                                                                                                                                                                       N
                                                                                                                                                                                       ACAAAAAACAAAAAATG 18
                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA sequencing; oligonucleotide; prinker; primer; linker; labelling region; cohesive end; complementary strand; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 94US-00275169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a
/note= "forms doubled stranded segment when bound to
nucleotides 5-22 of the sequence given in AAT12342"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
1. .18
                                                                                                                                                                                                                                                              0.3%;
                                                                                                                                                                                                                                            0,
                                                                                                                                                                                                                                                                   Score 15.4;
Pred. No. 8
                                                                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                Length 18;
                 variable region
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                                                                                                                                                                                                                                                 Gaps
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WPI; 1997-456759/42.
                                      Primi
                                                                                                                  09-NOV-1992;
18-OCT-1994;
                                                                                                                                                                            07-JUN-1995;
                                                                                                                                                                                                                   09-SEP-1997
                                                                                                                                                                                                                                                          US5665355-A
                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                          constant region;
                                                                                                                                                                                                                                                                                                                                                                         Antibody; T-cell receptor; beta chain; human immunodeficiency virus; HIV; blood; attenuation; primer; PCR; amplification; variable region;
                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                           (CONS-) CONSORZIO BIOTECNOLOGIE
                                                                                                                92US-00973485.
94US-00408011.
                                                                                                                                                                            95US-00488212
                                                                                                                                                                                                                                                                                                                                                      TCR; 88.
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The invention relates to a method for removing an antibody specific for TCR-V beta (T-cell receptor V beta protein) from an HIV-infected person, by removing blood from the person, removing the antibody from the blood, and reintroducing the blood into the person, thus allowing attenuation or aversion of immunodeficiency. The primers AAT98100-T98150 are used to check the efficiency of removal by detecting expression of the TCR-V-beta and V-alpha genes in a blood sample after treatment. This primer is targetted to the variable region sequence of the alpha chain gene and can be used in the amplification with primers AAT98148 or AAT98149 Removal of T-cell receptor-specific antibody from blood of HIV-infected person - by extracorporeal blood treatment, to attenuate or avert development of AIDS from HIV infection. Example 1; Col 11; 43pp; English.

Sequence 18 BP; 3 A; 7 C; 2 G; 6 T; 0 U; 0 Other;

Query Match Best Local (Matches 16; Similarity Conservative 94.1%; 0, Score 15.4; Pred. No. 8. Mismatches 6e+02 DB 1; Length 18; Indels 0; Gaps 0

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밁
                  Ś
                    4298 TTCGGAAGGAACTGGAG 4314
18
 TTCAGAAGGAACTGGAG 2
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RASSULT 688
AAXS5987/c
ID AAX859
XX AAX859
DT 13-SEP
XX PCR PT
XX ACQUIR
XW T Cell
KW CD4+ T
XX Synthe
OS Synthe
OS Homo 8
XX US5928
XX US5928
XX US5928
XX O9-NOV PCR primer used to amplify T cell receptor Va region cDNA. 13-SEP-1999 AAX85987 standard; DNA; 18 (first entry) ВP

Acquired immune deficiency syndrome; free antibody; paratope; epitope; T cell receptor variable beta region; TCR-V beta region; binding agent; CD4+ T cell; HIV; PCR primer; ss. 27-JUL-1999 18-OCT-1994; US5928642-A Homo sapiens Synthetic. 94US-00408011.

09-NOV-1992;

92US-00973485

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RESULT 689
AXX88163/c
ID AXX881
XX AXX881
XX D9-SEP
XX T cell
XX F cell
XX F cell
XX F CD4+ c
XX Synthe
XX US5925
XX US5925
XX O7-JUN
XX O9-NON
PR 09-NON
PR 18-OCT
XX Primi
XX Primi
XX Primi
XX WPI;
XX Primi
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XX Examp
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CC provi
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The specification describes a method for the diagnosis and treatment of acquired immune deficiency syndrome, in a person having free antibodies which have a paratope capable of binding to an epitope of a T cell receptor variable beta (TCR-V beta) region. The method comprises administering a binding agent homologous with the TCR-V beta epitope. The binding agent is useful in assays for detecting various CD4+ T cell subpopulations which carry particular V beta components. The binding agent is also useful in the treatment of people infected with HIV where it is able to remove an antibody able to bind with an epitope on a TCR-V beta cell in the blood of an infected person. The present PCR primer is used to amplify the TCR Va region, in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                            T cell receptor; beta chain; primer; antibody; paratope; AIDS; vaccine; epitope; TCR-V beta; immunogenic; anti-idiotypic; antiviral; detection; CD4+ cell subpopulation; acquired immune deficiency syndrome; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T cell receptor alpha chain primer V-alphal6.
This invention describes novel method for binding free antibodies having a paratope specific to an epitope on a T cell receptor (TCR-V beta) while providing an immunogenic substance able to raise anti-idotypic antibodies which bind to free antibodies bound at the same paratope
                                                                                                                                                                                                                                       09-NOV-1992;
18-OCT-1994;
                                                                                                                                                                                                                                                                                       07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-SEP-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX88163 standard;
                                                                                Example
                                                                                                                                            WPI; 1999-418267/35.
                                                                                                                                                                                                                                                                                                                      20-ЛИТ-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX88163;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-429481/36
                                                                                                                                                                                                          (CONS-) CONSORZIO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
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                                                                                <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTCAGAAGGAACTGGAG 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and treatment of acquired immune
                                                                                                             and treatment of acquired immune
                                                                                COT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                              11-12; 42pp;
                                                                                                                                                                                                                                       92US-00973485.
94US-00408011.
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                                                                                                                                                                                                                                                                                     95US-00488209
                                                                                                                                                                                                          BIOTECNOLOGIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         entry)
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                                                                            English.
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Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15.4; DB 1;
No. 8.6e+02;
                                                                                                             deficiency
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               deficiency syndrome.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                             syndrome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         specific to the epitope on the TCR-V beta and introducing this into a person to raise anti-idiotypic antibodies. The products of the inventic have antiviral activity and can be used in vaccines. The specific antibody binding affinities are useful in assays which detect the presence of CD4+ cell subpopulations carrying particular V beta components of the TCR-V beta in people infected with acquired immune deficiency syndrome (ALDS). AAX88119-X88169 represents primers used in the method of the invention
This is an antisense oligonucleotide targeted to the coding sequence, nucleotide 700, of human survivin DNA (see AAA08903). AAA08910-49 were analyzed for effect on survivin RNA levels by quantitative real-time PCR. The data obtained were averages from three experiments. This antisense oligonucleotide provided 18% inhibition of surivin MRNA. It was found that ISIS 23667 (AAA08925) provided 70% inhibition and ISIS 23672 (AAA08930) provided 64% inhibition. Survivin, an IAP (inhibitor of apoptosis) Caspase inhibitor, has been found to be involved in cell cycle regulation and is expressed in the 62/M phase of the cell-cycle in a cell cycle regulated manner and associates with microtubules of the mitotic spindle. Disruption of this interaction results in loss of survivin's anti-apoptotic function and increased caspase-3 activity during mitosis.
                                                                                                                                                                                                                                                                                                                                                             29-SEP-1998;
05-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-AUG-2000
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                                                                                                                                                                                            Example 15; Page 68; 73pp; English.
                                                                                                                                                                                                                           Antisense molecules targeted to in cancer cells.
                                                                                                                                                                                                                                                                                                     Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                          23-SEP-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200018781-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Survivin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human survivin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA08931;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA08931 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sapiens.
                                                                                                                                                                                                                                                                        2000-293103/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cycle regulation; cancer; cytostatic; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18 TTCAGAAGGAACTGGAG
                                                                                                                                                                                                                                                                                                                                   SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTCGGAAGGAACTGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inhibitor of apoptosis; IAP; caspase inhibitor; caspase-3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                     Ackermann EJ,
                                                                                                                                                                                                                                                                                                                                   PHARM INC
                                                                                                                                                                                                                                                                                                                                                               98US-00163162.
99US-00286407.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA antisense oligonucleotide, ISIS
                                                                                                                                                                                                                                                                                                                                                                                                          99WO-US022076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
1. .18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "phosphorothioate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
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                                                                                                                                                                                                                                                                                                    Swayze
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.4;
Pred. No. 8.
                                                                                                                                                                                                                                           Survivin,
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                                                                                                                                                                                                                                                                                                     Cowsert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    backbone"
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                                                                                                                                                                                                                                            useful
                                                                                                                                                                                                                                                                                                     3
                                                                                                                                                                                                                                           for inducing apoptosis
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RESULT 691
AAAS93/C
ID AAAS93
XX AAAS93
XX AAAS93
XX O1-NOV
XX NANOCI
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Best Loc
Matches
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Matches
                                                            Query Match
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                                                                                                                                                                        nonisotopic detection systems for blomolecules e.g. nucleic acids, proteins, lipids or drugs. The nanocrystals have polynucleotide strands attached to their surfaces with one end of the polynucleotide extending outwardly from the nanocrystal. The present sequence is one such polynucleotide. These nanocrystals are used with a second series of nanocrystals, which have polynucleotides complementary to the first polynucleotides, so that the respective complementary strands hybridise to each other and form a dendrimer. This dendrimer produces a signal which can then be detected e.g. fluorescence. The present sequence is composed mainly of TC repeats. This sequence may therefore be used with polynucleotide composed mainly of AG repeats (AAA58389)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Caspase-3 is associated with apoptotic cell death. It is therefore believed that survivin may counteract a default induction of apoptosis in the G2/M phase. It is also believed that the over expression of survivin in cancer may overcome this apoptotic check point, allowing undesired survival and division of cancer cells. Antisense oligonucleotides (ASO's) may be used to down regulate endogenous survivin and to increase caspase-3-dependent apoptosis in cells in the G2/M phase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Functionalized nanocrystal carrying polynucleotide, used for detecting target analyte, forms dendrimers with complementary nanocrystals to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Barbera-Guillem E, Nelson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-NOV-2000
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                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; Page 70; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amplify
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nanocrystal; biomolecule detection; nonisotopic detection system;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Polynucleotide # 6 used in a biomolecule detection system.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA58390;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA58390 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BIOC-) BIOCRYSTAL LTD
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hes 16;
                               Local
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the fluorescent signal.
                                                                                                                         18
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   Conservative
                                                                                                                         BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention relates to functionalised nanocrystals for use in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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99US-00437076
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
                            94.1%;
                                                                                                                         8 C; 1 G; 9 T; 0 U;
                                                            0.3%;
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0,
                                                            Score 15.4; DB 1; Length 18;
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Pred. No. 8.
                               Pred.
      Mismatches
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-hes 1; Indels
                               8.6e+02;
                                                                                                                         0 Other;
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   Indels
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Gaps
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ARASULT 692
ARAS8389
ID ARAS83
AC AR
RESULT 693
AAZ77145
ID AAZ771
XX
AC AAZ771
XX
DT 10-SEE
XX
DB Human
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to functionalised nanocrystals for use in nonisocopic detection systems for biomolecules e.g. nucleic acids, proteins, lipids or drugs. The nanocrystals have polynucleotide strands attached to their surfaces with one end of the polynucleotide extending outwardly from the nanocrystal. The present sequence is one such polynucleotide. These nanocrystals are used with a second series of nanocrystals, which have polynucleotides complementary to the first polynucleotides, so that the respective complementary strands hybridise to each other and form a dendrimer. This dendrimer produces a signal which can then be detected e.g. fluorescence. The present sequence is composed mainly of AG repeats. This sequence may therefore be used with polynucleotide composed mainly of TC repeats (AAA58390)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Functionalized nanocrystal carrying polynucleotide, used for detecting target analyte, forms dendrimers with complementary nanocrystals to amplify the fluorescent signal.
                                                                10-SEP-2001
                                                                                                                                                                               AAZ77145 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 18 BP; 8 A; 0 C; 10 G; 0 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 3; Page 70; 72pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Barbera-Guillem E,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-NOV-1998;
09-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200028088-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Polynucleotide # 5 used in a biomolecule detection system
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (BIOC-) BIOCRYSTAL LTD
                                                                                                                                                                                                                                                                                                                                                                                          1181 GAGAAAGAGAGAGAGAG 1197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1180
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Similarity 94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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                                                             (first entry)
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99US-00437076.
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                                                                                                                                                                                     DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.4;
Pred. No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8.6e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
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Human biallelic marker downstream amplification primer SEQ ID NO:11501.

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RESULT 694
AAZ72859
ID AAZ7288
XX AAZ7288
XX IO-SEE
DT 10-SEE
DT 10-SEE
DT Human
XX Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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Best Local &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ65654 to AAZ69578 represent human biallelic markers from the present invention, which contain a polymorphic base at position 24 of their nucleotide sequences. AAZ69579 to AAZ77440 represent amplification primers for the biallelic markers. The biallelic markers of the invention have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and dispostic methods, as well as the characterisation of the differential efficacious responses to and side effects from the observation of the development of the differential efficacious responses to and side effects from the development of the differential efficacious responses to and side effects from the development of the differential efficacious responses to and side effects from the development of the differential efficacious responses to and side effects from the development of the differential efficacious responses to and side effects from the development of the differential efficacious responses to and side effects from the development of the differential efficacious responses to and side effects from the development of the development of
                                                                                Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; bolymorphic base; genotyping; haplotyping; high didation; identification; characterisation; haplotyping; single nucleotide polymorphism; SNP; PCR primer; amplification; single nucleotide polymorphism;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotrype; polymotrphic base; genotyping; haplotyping; hybridisation; identification; characterisation; haplotyping; hybridisation; dentification; SNP; PCR primer; amplification; single nucleotide polymorphism; SNP; PCR primer;
                                                                                                                                                                                                                         Human
                                                                                                                                                                                                                                                                            10-SEP-2001
                                                                                                                                                                                                                                                                                                                                     AAZ72859
                                                                                                                                                                                                                                                                                                                                                                                      AAZ72859 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 18 BP; 7 A; 0 C; 9 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pharmaceutical agents acting on a disease as well as other treatment. N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-013267/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cohen
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23-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO9954500-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel biallelic markers used to construct a high density disequilibrium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-OCT-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2436 GGATGAGAAGGGGAGAG 2452
                                                                                                                                                                                                                      biallelic marker upstream amplification primer SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGATAAGAAGGGGAGAG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       98US-0082614P
98US-0109732P
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                                                                                                                                                                                                                                                                                                                                                                                         DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                         BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.4;
Pred. No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 18; .6e+02;
                                                                                                                                                                                                                         ID NO:7215.
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RESULT 695
AAS21649
ID AAS216
AC AAS216
AC AAS216
XX 21-NOV
XX 21-NOV
XX Surviv
KW Surviv
KW hyperr
XX Hymon 6
OS Hymon 6
COS Synthe
XX W02001
XX W02001
XX W02001
XX W02001
XX W02001
XX W02-FEE
XX W1-FEE
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       have a variety of uses: they can be used for high density mapping of the human genome, and in complex association studies and haplotyping studies which are useful in determining the genetic basis for disease states. Compositions and methods of the invention can also be useful for the identification of the targets for the development of pharmaceutical agents and diagnostic methods, as well as the characterisation of the differential efficacious responses to and side effects from pharmaceutical agents acting on a disease as well as other treatment.

N.B. The SEQ ID NOs 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not actually given a sequence in the Sequence Listing from the
                                                                                                                                                                                                                                                 Human Survivin antisense oligonucleotide #114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAZ65654 to AAZ69578 represent human biallelic markers from the pres
invention, which contain a polymorphic base at position 24 of their
nucleotide sequences. AAZ69579 to AAZ77440 represent amplification
primers for the biallelic markers. The biallelic markers of the inve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-013267/01.
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23-NOV-1998;
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                              02-FEB-2000; 2000US-00496694.
                                                                                           09-AUG-2001
                                                                                                                          WO200157059-A1
                                                                                                                                                       Synthetic.
                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                  hyperproliferative
                                                                                                                                                                                                                   Survivin;
                                                                                                                                                                                                                                                                                21-NOV-2001
                                                                                                                                                                                                                                                                                                              AAS21649;
                                                                                                                                                                                                                                                                                                                                           AAS21649 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel biallelic markers used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-APR-1999;
   (ISIS-) ISIS PHARM INC
                                                           30-JAN-2001; 2001WO-US002939.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GEST ) GENSET.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    4871
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                                                                                                                                                                                                                                                                                                                                                                                                                                         CTCAGTTTCTTTCTCTG 4887
                                                                                                                                                                                                                   human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Blumenfeld
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 1 A; 6 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98US-0109732P.
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                                                                                                                                                                                                                    mouse;
                                                                                                                                                                                                      ouse; cytostatic; antisense oligonucleotide;
condition; cancer; apoptosis; cytokinesis; ss.
                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4; DB 1
Pred. No. 8.6e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            construct a high density disequilibrium
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 18
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RESULT 696
AAS21598
ID AAS215
XX
AC AAS215
XX
DT 21-NOV
XX
Human
XX
Synthe
XX
OS Homo (
OS Homo (
OS Homo (
OS Homo (
OS Synthe
XX
PN WO200:
XX
PN WO200:
XX
PF 30-AU
XX
PF 30-FE
XX
PF 30-FE
XX
PF 30-FE
XX
PF WPI;
XX
PF WPI;
XX
PF Novel
PT treat
XX
PS Examy
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to antisense oligonucleotides targeted to acid molecule encoding human Survivin, where the antisense oligonucleotide inhibits the expression of human Survivin. These
                                                                                                                                                                                                                                                                                                                                             Survivin; human; mouse; cytostatic; antisense oligonucleotide; hyperproliferative condition; cancer; apoptosis; cytokinesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAS21598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 18
   Example 16; Page 54; 120pp; English
                                                Novel antisense compounds
                                                                                                                                                                          02-FEB-2000; 2000US-00496694
                                                                                                                                                                                                        30-JAN-2001; 2001WO-US002939.
                                                                                                                                                                                                                                       09-AUG-2001.
                                                                                                                                                                                                                                                                    WO200157059-A1
                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                  Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                             21-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAS21598 standard; DNA; 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            treatment of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1184 AAAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                                                                            Survivin antisense oligonucleotide #64.
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                                                                                                            CF,
                                                                                                                                           SISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 11 A; 0 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                            Ackermann
                                                                                                                                           PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                for modulating
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                                                                                                              Swayze
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15.4; DB 1; Length 18; Pred. No. 8.6e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                              器,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BB,
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                                                                                                              Cowsert
                                                  the
                                                  expression of Survivin
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RESULT 697
AMS21558
ID AMS215
AC AMS215
XX AMS215
XX AMS215
XX Surviv
DH Claman
XX Surviv
KW Synthe
XX Homo &
OS Synthe
XX O2-FEF
XX O2-FEF
XX O2-FEF
XX O2-FEF
XX USIS-
PH Bennet
XX PF 30-JAN
XX PF 30-JAN
XX PF 30-JAN
XX PF 10vel
PH Novel
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PH Novel
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CC antis
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Best Local
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acid molecule encoding human Survivin, where the antisense oligonucleotide inhibits the expression of human Survivin. These antisense oligonucleotides are used in the treatment of an animal suffering from a disease or condition associated with Survivin, e.g. a hyperproliferative condition such as cancer, and comprises administering a therapeutically or prophylactically effective amount of the antisense oligonucleotide so that expression of Survivin is inhibited. The
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-JAN-2001; 2001WO-US002939
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Survivin; human; mouse; cytostatic; antisense oligonucleotide; hyperproliferative condition; cancer; apoptosis; cytokinesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           21-NOV-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Survivin, used in the method of the invention
                                                                                                                                                                    The invention relates to antisense oligonucleotides targeted to a
                                                                                                                                                                                                                      Example 15;
                                                                                                                                                                                                                                                                  treatment of cancer
                                                                                                                                                                                                                                                                                         Novel antisense compounds for
                                                                                                                                                                                                                                                                                                                                           WPI; 2001-488863/53
                                                                                                                                                                                                                                                                                                                                                                                       Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-FEB-2000; 2000US-00496694
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human Survivin antisense oligonucleotide #24.
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16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                       ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAAGAGAGAGAGAGA 17
                                                                                                                                                                                                                   Page 53; 120pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                       Ackermann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                       Swayze
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.4;
Pred. No. 8.
                                                                                                                                                                                                                                                                                         modulating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                       88,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 B.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                          Cowsert
                                                                                                                                                                                                                                                                                              the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 18;
                                                                                                                                                                                                                                                                                           expression of Survivin
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Matches 16
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                         This invention describes a novel method of analysing an HIV target nucleic acid which comprises amplifying the target sequence using primers specific for the HIV pol region (encoding the viral protease and reverse transcriptase) and analysing the amplified product. The invention is used to determine viral genotype of HIV infections by detecting known drug resistance mutations in the HIV pol gene. The detection of mutations conferring drug resistance in the HIV pol gene is significant in determining drug sensitivity of the virus. A 1.57 kilobase region of the HIV pol gene is a particularly important region containing clinically
                                                                                                                                                                                                                                                                                                                                                                                                  Amplifying the HIV pol region using specific primers and a sequencing the product using specific sequence primers allows detection of known drug resistance mutations and so determination of viral genotype.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-506465/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Survivin, used in the method of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-SEP-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HIV Pol sequencing primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-SEP-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABL61267 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Johnston-Dow LA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (/NHOL)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               gene; prime infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1184 AAAGAGAGAGAGAAA 1200
                                                                                                                                                                                                                                                                                                                                            8; Col 13-14; 12pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DEMETER L. WHITE C B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MYERS A.
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CONRAD M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JOHNSTON-DOW L
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         mutation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Myers A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               viral protease; reverse transcriptase; genotyping;
tection; drug resistance mutation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Demeter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA;
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   sequence
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   segencing
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ABS97214 standard;
                                                                                                                                                                                                                                       Example 3; Page 103; 714pp; English
                                                                                                                                                                                                                                                                                        disorder-related traits.
                                                                                                                                                                                                                                                                                                         Isolated nucleic acid molecules having polymorphisms in known human
e.g. cytochrome p450 and cathepsin S useful as genetic linkage marke
for locating, identifying and characterizing the genes responsible i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-NOV-2000; 2000US-00724389.
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Pred. No. 8.
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linkage markers
responsible for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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This invention relates to the sequence of an isolated nucleic acid molecule comprising at least one base variation from that of a known human cytochrome P450 A2 (CYP4501A1), cytochrome P450 A2 (CYP4501A2), cytochrome P450 D22 (CYP45002E1), adrenergic receptor betal (ADBR1), aryl hydrocarbon (AHR), aryl hydrocarbon receptor nuclear translocator (ARNT), cathepsin S (CTSS), cyclooxgenase 2 (COX2), diazepam binding inhibitor (DBI), epoxide hydroxylase 2 (EPHX2), 5-lipoxygenase activating protein (FLAP), glutathione-S-transferase 12 (GST12), histamine-N-methyl transferase (HNMT), (kallikrein 2) KLK2, nicotinamide -N-methyl

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RESULT 700
ABI82302/c
ID ABI823
XX Human;
XX Infect
XX I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC transferase (NNMT), NADPH quinone oxidoreductase 2 (NQO2),
CC (UGT2B4), UDP-glucuronosyl transferase 2B4
CC (UGT2B4), UDP-glucuronosyl transferase 2B7
CC (MDR1), lactotransferrin (LTF), multidrug resistance resistance 1
CC (MDR1), lactotransferrin (LTF), multidrug resistance associated protein 3
CC (MDR1), lactotransferrin (LTF), multidrug resistance associated protein 3
CC (MRP3), orphan nuclear receptor (NR112), or actylcholine muscarinic
CC receptor 1, 2, 3, 4, or 5 (CMMR1, CMMR2, CMMR4 or CTMR5) sequence.
CC The polymorphisms in the human genes cited in the invention are useful as
CC genetic linkage markers for locating and characterising the genes that
CC are responsible for specific traits within the genome and eventually
CC identifying the genes responsible for a variety of disorder-related
CC expression, mutation or underexpression, which may be used in diagnosing
CC and/or treating the disorders. The nucleic acid molecules comprising the
CC polymorphic sequences contained in CYP4501A1, CYP94501A2, CYP4502E1,
CC ARMT, EPHX2, GST12, NNMT, NQO2, NR112, STM, UGT2B4, UGT2B7, UGT2B15, AHR,
CC MDR1 and/or MDR3 are useful for screening individuals for altered drug
CC ausceptibility to cancer. Polymorphic sequences in ADRB1 or CHMR2 are
CC used to screen for altered cardiovascular function, in COX2 for altered
CC used to screen for altered cardiovascular function, in COX2 for altered central
CC immunological function, in FLAP and HNMT for altered guimonary,
CC immunological function, in CHMR3, CHMR4 or CHMR5 for altered central and
CC peripheral nervous system function. The present sequence represents a
CC sequencing primer used to sequence the polymorphic genes of the invention
CC sequencing primer used to sequence the polymorphic genes of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 p53 mutation
                                                                                                                                                                                                                                                                                                                                                                                                   04-APR-2001; 2001WO-US010958.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO200179548-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABI82302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABI82302 standard; DNA; 18 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18
                                                                                                                               Barany F,
                                                                                                                                                                                                                                                                                                            14-APR-2000; 2000US-0197271P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4420 CTGGTGTGGGAGGCCTT 4436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17
                                                                                                                                                                                                                          CORNELL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTGGTGTGGGCGGCCTT 1
                                                                                                                               Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            detection primer/probe #181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                          RES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; 9 C; 4 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.3%;
94.1%;
                                                                                                                                                                                                                      FOUND INC
                                                                                                                                        Gerry NP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.4;
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                    Favis R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 18;
                                                                                                                                        Kliman
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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문 S

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cc e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal infectious agents e.g. Cryptococcus neoformans, Candida albicans and Cappergillus fumigautus, viruses e.g. T-cell lymphocytotrophis cirus, Epstein-Barr virus and polio virus, and parasitic infectious agents concluded agents of the conclusion of the concerning cancer involving oncogenes, tumour suppressor genes, or genes involved in DNA amplification, replication, recombination or repair, the cancer is specifically associated with a gene selected from BRCA1 gene, p53 gene, human papillomavirus types 16 and 18 and liver cancers. The method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning collectron microscope and infrared microscope) the support at the particular sites and identifying if ligation to a presence or absence of the target nucleotide sequences. ABI82074 to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes a method (M1) for designing capture oligonucleotide probes (I) for use on a support to which complementary oligonucleotide probes (II) will hybridise with little mismatch, where (I) have melting temperatures within a narrow range. The method is useful for detecting infectious diseases caused by bacterial infectious agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               300pp; English.
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Query Match Best Local : Matches 15-FEB-2002 ABI82323; ABI82323 standard; DNA; 18 4737 GGAGACCCATCCTCACC 4753 16; Similarity Conservative (first entry) 94.18; ВP 0; Mismatches Score 15.4; Pred. No. 8 8.6e+02; Indels 0; Gaps

밁 8

Sequence 18 BP; 3 A; 4 C; 8 G; 3 T; 0 U; 0 Other;

0.3%;

DB 1;

Length 18;

0

the present

invention

Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss. Synthetic Homo sapiens p53 mutation detection primer/probe #202.

RESULT 701
AB18232/c
ID AB1823
XX AB1823
XX AB1823
XX AB1823
XX AB1823
XX IS-PEB
XX P53 mu
XX Human;
KW Human;
KW Infect
KW Oncoge
KW On WPI; 2002-034366/04 Barany F, 14-APR-2000; 2000US-0197271P 04-APR-2001; 2001WO-US010958 25-OCT-2001 WO200179548-A2 (CORR) CORNELL RES Zirvi M, FOUND INC Gerry NP, Favis Kliman

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RESULT 702
ABI82303/c
ID ABI823
XX ABI823
XX ABI823
XX ABI823
XX IS-FEB
XX I
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a method (M1) for designing capture CC oligonuclectide probes (II) will hybridise with little mismatch, where CC (I) have melting temperatures within a narrow range. The method is useful CC (I) have melting temperatures within a narrow range. The method is useful CC e.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal CC infectious agents e.g. Cryptococcus meoformans, Candida albicans and CC Aspergillus fundagautus, viruses e.g. T-cell lymphocytotrophis cirus, Esstein-Barr virus and polio virus, and parasitic infectious agents CC medinesis. The method is also useful for detecting genetic diseases such CC as 21 hydroxylase deficiency, Turner Syndrome and obesity defects. CC Detecting cancer involving oncogenes, tumour suppressor genes, or genes CC involved in DNA amplification, replication, recombination or repair, the CC cancer is specifically associated with a gene selected from BRCA1 gene, human papillomavirus types 16 and 18 and liver cancers. The CC grant for also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning cleature sites and infrared microscope) the support at the CC generace or absence of the target nucleotide sequences. AB182074 to a feel for the present onligonucleotide sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; K-ras; PCR primer; probe; capture prove, municipal disease; ligase detection reaction; LDR; p53; BRCA1; BRCA2; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndrome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; oncogene; tumour suppressor; human papillomavirus; forensic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABI82303 standard; DNA; 18
                                         Barany F,
                                                                                                                                                                          14-APR-2000; 2000US-0197271P.
                                                                                                                                                                                                                                        04-APR-2001; 2001WO-US010958
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 p53 mutation detection primer/probe #182.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 3; Page 67; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
                                                                                                                                                                                                                                                                                                             25-OCT-2001
                                                                                                                                                                                                                                                                                                                                                                            WO200179548-A2
                                                                                                           (CORR ) CORNELL RES FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4737 GGAGACCCATCCTCACC 4753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGAGGCCCATCCTCACC 1
                                         Zirvi M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BP; 2 A; 4 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
                                         Gerry NP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                         Favis R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15.4;
No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1; Length 18; .6e+02;
                                             Kliman R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                  Pyrin domain; PYD domain; antiinflammatory; antiparkinsonian; antiarteriosclerotic; antipsoriatic; antibacterial; virucide; neuroprotective; antiarthritic; antirheumatic; nephrotropic; osteopathic; nootropic; intracellular signal transduction; inflammation; Alzheimer's disease; infection; psoriasis; asthma; arteriosclerosis; multiple sclerosis; rheumatoid arthritis; sarcoidosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Designing capture oligonucleotide probes for use on a support to which complementary oligonucleotides hybridize with little mismatch.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2002-034366/04.
                                     15-NOV-2000; 2000DE-01056687.
30-NOV-2000; 2000DE-01059595.
                                                                                                                                                                                                                                                                                                                                                                                                     Pyrin domain containing protein coding sequence PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 18
                                                                                             30-OCT-2001; 2001WO-EP012545.
                                                                                                                                                                         WO200240668-A2
                                                                                                                                                                                                            Unidentified.
                                                                                                                                                                                                                                                  osteoarthritis; glomerulonephritis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                             20-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL47145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAL47145 standard; DNA; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 3; Page 66; 300pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present invention describes a method (M1) for designing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4737 GGAGACCCATCCTCACC 4753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGAGGCCCATCCTCACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 3 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.4; DB 1; Pred. No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         capture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                <u>.</u>,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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RESULT 704
ADB99917
ID ADB9999
XX ADB9999
XX ADB999
XX ADB999
XX ADB999
XX OA-DEC
XX LRP5 II
XX OSteep
KW Done II
XX Done II
XX II-MAY
PR 11-MAY
P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 16
                                                                                                       New nucleic acid comprising a mutation in LRP5 or LRP6, useful diagnosing a HBM-like phenotype in a subject and for preparing composition for modulating bone mass and/or lipid levels in a suffering from e.g. osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                  11-MAY-2001; 2001US-0290071P.
17-MAY-2001; 2001US-0291311P.
01-FBB-2002; 2002US-0353058P.
04-MAR-2002; 2002US-0361293P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Osteopathic; Gene therapy; High Bone Mass; HBM; LRP5; Zmax1; LRP6; bone mass modulation; osteoporosis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to treat diseases associated with impaired intracellular signal transduction, particularly inflammation such as psoriasis, arteriosclerosis, bacterial or viral infections (particularly meningitis and pneumonia), multiple sclerosis, rheumatoid arthritis, asthma, sarcoidosis, glomerulonephritis and osteoarthritis, and also Alzheimer's and Parkinson's diseases. The present sequence is a PCR primer used to isolate a coding sequence of the invention
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                                                          Disclosure;
                                                                                                                                                                                                                                                WPI; 2003-129214/12.
                                                                                                                                                                                                                                                                                                                                                       (GENO-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tachopp J,
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                                                       Page 53; 629pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 4 A; 3 C; 8
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                                                                                                                                                                                                                                                                                                     Yaworsky PJ,
                                                                                                                                      a subject
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present invention relates to High Bone Mass (HBM), LRP5 (Zmax1) and

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Best Local S
Matches 16
Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                         The present invention relates to a method for amplifying and sequencing HIV pol region DNA. Also disclosed is a kit comprising first and second primers, and a temperature-stable DNA polymerase. The invention also discloses a method for obtaining de novo sequence information for different HIV quasi-species present in a patient's sample. The kit is useful for HIV sequencing and genotyping, particularly for amplifying regions of the HIV pol gene. It is also useful for automated sequence analysis and determination of heterozygosity. The present sequence represents a sequencing primer for HIV pol region DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New kit comprising first and second primers, and a temperature-stable DNA polymerase, useful for HIV sequencing and genotyping, and for automated sequence analysis and determination of heterozygosity.
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                                                                                                          Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 3; SEQ ID NO 5; 11pp; English.
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Conrad M, Myers A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA amplification; sequencing; HIV pol; genotyping; sequence analysis; heterozygosity; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADI34074 standard; DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (APPL-) APPLERA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3422 TGAGGCAGGAACTGAGG 3438
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                                                                                                                9 C; 3 G; 0 T; 0 U; 0 Other;
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Pred. No. 8.6e+02;
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     Mismatches
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Best Local Similarity
Matches 16; Conserv
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       06-MAY-2004
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                                                  ADI20873;
                                                                                               ADI20873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 8; SEQ ID NO 5; 14pp; English.
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Conrad M, Myers A;
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05-MAR-2002; 2002US-00092022.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequencing primer #1 for HIV pol gene
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WHITE C B.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JOHNSTON-DOW L A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        KOHLENBERGER R. CONRAD M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SONG K.
                                                                                                                                                                                                                                                     AGCCTACAGCCCCACCA 1277
                                                                                               standard;
                                                                                                                                                                                                            AGCCAACAGCCCCACCA 17
                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first
    (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Demeter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
                                                                                            DNA; 18
                                                                                                                                                                                                                                                                                                                        0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
    entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                          invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18
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                                                                                                 ΒP
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                                                                                                                                                                                                                                                                                                                        Score
Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        White CB,
                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                        15.4; DB 1; Length 18; No. 8.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Song K,
                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kohlenberger
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                                                                                                                                                                                                                                                                                                     Gaps
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RESULT 708
ADQ88986/c
ID ADQ889
XX
AC ADQ889
XX
AC ADQ889
XX
DT 23-SEP
XX
DE Breast
XX
                                                                                                                                                                                                                                                                                                                                                                             The present invention relates to detecting the methylation state of the CC s' and promoter region of the gene DD3 within a subject comprising CC contacting a target nucleic acid having one or more sequences selected CC from 5 3581 base pair sequences in a biological sample with at least one reagent or a series of reagents. The method is useful for detecting the CC subject. The set of oligonucleotides comprising at least three of the CC oligoners is useful for detecting the cytosine methylation state and/or Single nucleotide polymorphisms (SNPs) within SRO. ID NO. 1-5 and its complementary sequences. The set of oligoners is also useful for detecting the methylation state of all CpG dinucleotides within SRO ID NO. 1 and its complementary sequences. The set of oligonucleotides within SRO ID NO. 1 and its complementary sequences. The set of all CpG dinucleotides within SRO ID NO. 1 and its complementary sequences. The set of all CpG dinucleotides for the CC amplification of DNA sequences selected from SRO ID NO. 1-5 and its comprising at least one mucleic acid. The methylation state of the gene DD3 CC comprising diseases associated with the methylation state of the gene DD3 CC oligonucleotide or PNA-oligomer, kit, array or the set of oligonucleotides is useful for the characterization, classification, CC differentiation, grading, staging, and/or diagnosis of cell proliferative disorders. It can also be used for the therapy of cell proliferative disorders. The present sequence represents a detection oligonucleotide of the invention.
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                                                                                                                                                                                                                                                                                                   Best Loc.
Matches
                                                                                                                                                                                                                                                                                                                                      Query Match
                     Breast cancer
                                                                                                                                                                                                                                                                                                                                                                             Sequence 18 BP; 2 A; 0 C; 1 G; 15 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Detecting methylation of 5' and promoter region of DD3 gene for diagnosing proliferative disorders comprising contacting target nucleic acid with a reagent that distinguishes between methylated and non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MS SnuPE detection oligonucleotide for
                                                        23-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          methylated CpG dinucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-JUL-2002; 2002DE-01030692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-JAN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO2004005543-A1
                                                                                             ADQ88986
                                                                                                                                  ADQ88986 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (EPIG-) EPIGENOMICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JUN-2003; 2003WO-EP006690
                                                                                                                                                                                                                                                                                                                     Local
                                                                                                                                                                                                                                                    5389 AATTAAAAAAATACAAA 5405
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2004-091385/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CpG dinucleotide; cell proliferative disorder; ss.
                                                                                                                                                                                                                             17
                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 76; 56pp; English.
                                                                                                                                                                                                                                                                                                     Conservative
                                                        (first entry)
                 associated polymorphism detection
                                                                                                                                  DNA; 18
                                                                                                                                                                                                                                                                                                                      94.1%;
                                                                                                                                  ВP
                                                                                                                                                                                                                                                                                                   0,
                                                                                                                                                                                                                                                                                                                        Score 15.4;
Pred. No. 8.
                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DD3
                                                                                                                                                                                                                                                                                                                        8.6e+02;
                                                                                                                                                                                                                                                                                                                                        DB 1; Length 18,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            #13
                     primer #84.
                                                                                                                                                                                                                                                                                                     0
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0;

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RESULT 709
ANTO9139
ID ANTO91
XX ANTO91
XX AOTO91
XX O2-AUG
DT 02-AUG
XX Suppre
KW Suppre
KW Suppre
KW immune
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX W0953!
XX W0953!
XX PD 28-DE(
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention describes a method of identifying (M1) a subject at risk of breast cancer. The method comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject. The methods, nucleic acids, proteins, and compositions are useful for diagnosing, preventing, and treating breast cancer. Also described is a method useful for identifying candidate therapeutics for treating breast cancer. This sequence represents an extension primer used to analyse polymorphisms associated with breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Identifying a subject at risk of breast cancer comprises detecting the presence or absence of one or more polymorphic variations associated with breast cancer in a nucleic acid sample from a subject.
                                                                                                                                                                  Synthetic
                                                                                                                                                                                            Suppression; nuclear factor kappa-B; primer; septic shock; cytokine; immune response; retrovirus; infection; HIV; antisense; translation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 2; Page 61; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-NOV-2002; 2002US-0429136P
24-JUL-2003; 2003US-0490234P
  28-DEC-1995
                                                                                misc_feature
                                                                                                                        misc_feature
                                                                                                                                                                                                                                                                 02-AUG-1996
                                                                                                                                                                                                                                                                                                                       AAT09139 standard; DNA; 19 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Roth RB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-NOV-2003; 2003WO-US037831.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUL-2004.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cytostatic; gene therapy; breast cancer; polymorphism detection; PCR; primer; ss; extension primer.
                            WO9535032-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO2004055196-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (SEQU-) SEQUENOM INC
                                                                                                                                                                                                                                                                                                                                                                                                                   3899 AGATTGAATTCTGTGTG 3915
                                                                                                                                                                                                                                                                                                                                                                                            17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nelson MR,
                                                                                                                                                                                                                                      construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 6 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                (first entry)
                                                                                    18. .19
                                                                                                                          Location/Qualifiers
                                                        note= "internucleotide phosphothicate linkage"
                                                                                             /*tag= b
/note= "internucleotide
                                                                     *tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                                                     sense primer binds bases 718-737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Braun A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     μ
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Pred. No. 8.6e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             4; DB 1; Length 18; 8.6e+02;
                                                                                               phosphothicate linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Reneland
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     77
                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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RESULT 710
AATO9138/c
ID AATO91
XX AATO91
XX O2-AUG
DT 02-AUG
XX Suppre
KW Suppre
KW Suppre
KW Suppre
KW Synthe
FT misc_f
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The oligonucleotide AAT09138-40 were used in an assay to test for inhibition of translation of the tax protein and transactivation of the HTLV-1 long terminal repeat (LTR) in a HTLV-1 LTR/tax construct in mouse fibroblastic tumours generated in transgenic mice overexpressing the tax protein. The oligonucleotides acts to suppress tax protein translation which cannot then activate the HTLV-1 LTR and leading to a suppression of potential tumourogenic growth of cells. This sense oligonucleotide corresp. to bases 718-737 of an HTLV-1 LTR/tax construct (see Nerenberg et al., Science, 237: 1324 (1987). Similarly a novel method of suppressing nuclear factor kappa-B (NF-kB) dependent processes such as septic shock or disorders mediated by immune or cytokine responses or retroviral infections, partic. HIV, comprises novel antisense oligonucleotides (AAT09134-7) targeted to translational start site sequences (AAT09130-3) of NF-kB such that translation from the start site
                                                                                                                                                                                                                                              misc_feature
                                                                                                                                                                                                                                                                                                                                   misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Suppression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HTLV-1/tax construct antisense primer binds bases 718-737.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-AUG-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT09138;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAT09138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19 BP; 3 A; 9 C; 2 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     on the NF-kB mRNA is prevented
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 1; Page 15; 52pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New anti:sense oligo:nucleotide(s) against nuclear factor-kappa B subunit mRNA - used to suppress NF-kB dependent processes in individuals, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1996-058150/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-AUG-1994;
                                                                                                                                                                    misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     septic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nerenberg MI,
                                                              WO9535032-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (SCRI ) SCRIPPS RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4066 TTCCARATGGCCCACTT 4082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   l Similarity
16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         response;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTCCACATGGCCCACTT 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nuclear factor kappa-B; primer;
nse; retrovirus; infection; HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kitajima I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9308-00110161.
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                                                                                                                                                                                                                                              /*tag= a
/note= "translational initiation codon'
17. .18
                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                 note= "internucleotide
                                                                                                                                                                                           note= "internucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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Pred. No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                 phosphothicate linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; DB 1;
B.7e+02;
                                                                                                                                                                                              phosphothicate linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense; translation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  septic shock; cytokine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 19
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28-DEC-1995

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RESULT 711
AAT46087
ID AAT460
XX AAT460
XX Primex
XX 19-FEE
XX STS; E
KW Yeast
XX Synthe
XX Synthe
XX 17-SEI
XX 17-SEI
XX 17-API
XX 10-IM

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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The oligonucleotide primers AAT09138-40 were used in an assay to test for inhibition of translation of the tax protein and transactivation of the HTLV-1 long terminal repeat (LTR) in a HTLV-1 LTR/tax construct in mouse fibroblastic tumours generated in transgenic mice overexpressing the tax protein. The oligonucleotides acts to suppress tax protein translation which cannot then activate the HTLV-1 LTR and leading to a suppression of potential tumourogenic growth of cells. This antisense oligonucleotide is taxgeted to bases 718-737 of an HTLV-1 LTR/tax construct (see Nerenberg et al., Science, 237: 1324 (1987). Similarly a novel method of suppressing nuclear factor kappa-B (Nr-KB) dependent processes such as septic shock or disorders mediated by immune or cytokine responses or retroviral infections, partic. HIV, comprises novel antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      retroviral infections, partic. HIV, comprises novel antisense oligonucleotides (AAT09134-7) targeted to translational start site sequences (AAT09130-3) of NF-kB such that translation from the starts.
                               WPI; 1996-432990/43
                                                                                                                                                                                                                                                                                                                                                              US5556786-A
                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Primer for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         19-FEB-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19
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                                                                                                                                                                                                                                                   27-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 1; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New anti:sense oligo:nucleotide(s) against nuclear factor-kappa B subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20-AUG-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-AUG-1994;
                                                                                                                                                                                                                                                                                                        17-SEP-1996
                                                                                                                                          (UNIW ) UNIV WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (SCRI ) SCRIPPS
                                                                                 La Chapelle A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4066 TTCCAAATGGCCCACTT 4082
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence-tagged site; primer; EDA; anhidrotic ectodermal dysplasia;
t artificial chromosome; Homo sapiens; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NF-kB mRNA is prevented
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STS associated with EDA gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              to suppress NF-kB dependent processes in individuals,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry
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                                                                                                                                                                                             93US-00052997
                                                                                                                                                                                                                                                   93US-00052997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; 2 C; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                 Kere J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  52pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G; 3 T; 0 U; 0 Other;
                                                                                    Schlessinger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ŏ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 19;
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FFX8X25555555555
                                                                       artificial chromosome (YAC) clones from the region using RFLP markers and new unique markers. AAT46079-92 are primers for sequence-tagged sites associated with the anhidrotic ectodermal dysplasia (EDA) gene. AAT46083-92 are associated with new markers labelled sWXD632, sWXD178, sWXD634,
                                                                                                                                                                                                                                                                                                                                                                                           Cloning vector contg. the human anhydrotic ecto-dermal dysplasia gene for diagnosis of EDA related diseases.
                                                                                                                                                                                                                                                                                                                                    Claim 5; Col 28; 19pp; English.
                                                                                                                                                                              hypoplasia of hair, teeth and sweat glands. The EDA gene has been mapped to Xq12-q13 by genetic linkage analysis using restriction fragment length polymorphisms (RFLP) markers. Translocation breakpoints were also used to golymorphisms (RFLP) markers as well as the recovery of yeast
                                                                                                                                                                                                                                                                                         an X-chromosomal recessive disorder linked with the absence or
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Query Match
Best Local Similarity
Matches 16; Conserv
                339
N
TTTCCTACCACTCCACC 18
                TTTCCTACCACTCCCC 355
                                   Conservative
                                            0.3%;
                                   0,
                                            Score 15.4;
Pred. No. 8.
                                     Mismatches
                                           8.7e+02;
                                                     DB 1;
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                                     Indels
                                                      19
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                                     Gaps
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Sequence 19 BP; 4 A; 10 C; 0

G; 5 T; 0 U; 0 Other;

sWXD635 and sWXD636

F S

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RESULT 712
AAVO1290/c
ID AAVO12
XX AAVO12
XX AAVO12
XX PCR pr
KW PCR pr
KW Univer
XX Synthe
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-AUG-1997
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primer; polymerase chain reaction; amplification; UM-STS; universal mammalian sequence tagged site; genomic map; clone;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-MAR-1998
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New oligonucleotide primers amplifying gene regions conserved among mammals - useful for developing genomic maps, isolating clones and Claim 1; Page 11; 26pp; English. cross-species comparisons. and making

WPI; 1997-435083/40. Brewer GJ, Venta PJ, (UNMI)

UNIV MICHIGAN.
UNIV MICHIGAN STATE.

Yuzbasiyan-Gurkan

The present sequence represents a specifically claimed oligonucleotide PCR primer. The oligonucleotide can be used for polymerase chain reaction (PCR) amplification of DNA, specifically regions of specific genes that are conserved among mammalian species, i.e. pairs of oligonucleotides from the present specification represent universal mammalian sequence-tagged site (UM-STS) primers. The primers are used to develop genomic maps, to isolate clones from libraries, to make cross-species comparisons and to develop additional genetic markers. UM-STS allow genomic comparisons to be made between more вресіев

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RESULT 713
ANT63656/c
ID ANT636
XX ANT636
XX ANT636
XX Oligo
DE Oligo
XX antise
XX ANT636
XX Oligo
XX Oligo
XX OP-AUG
XX OP-AU
                                                                                                         RESULT 714
AAV05874
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                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotides having a partial sequence consisting of at least 15 bases of AATG3641 (an antisense oligo complementary to a region of the tax gene which can inhibit human T-cell lymphotropic virus type 1 (HTIV-1) viral antigen expression) are claimed. In an example, six antisense oligos were designed, T1-T6 (AATG3650-55) and were compared to six oligos derived from other regions of HTLV-1, i.e. SJI (splice junction), P1 (p21), R1 (rex), RR1 (rex response element), E1 (env) and G1 (gag), four reference oligonucleotides TIS (tax-sense), HC (dC20), HT (dT20) (AATG3647-49) and a random 20mer (RAN) in a HTLV-1 virus antigen expression inhibiting test. Oligonucleotide T1 gave the best results. The present sequence is an oligonucleotide disclosed in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anti-HTIV-1 anti-sense oligo:nucleotide - is complementary to region of tax gene from human T-cell lymphotropic virus type 1 and inhibits viral antigen expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 5 A; 2 C; 9 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense; complementary; tax gene; inhibit; HTLV-1;
human T-cell lymphotropic virus type 1; viral antigen expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligo disclosed in patent about Anti-HTLV antisense therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-JUN-1997
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                          AAV05874;
                                                                                 AAV05874 standard; DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure, Page 9; 10pp; Japanese
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (SOYA-) SOYAKU GIJUTSU KENKYUSHO
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                                                                                                                                                                                                                                                                                                                                   16;
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                                                                                                                                                                                                                                                                                                                                                                Similarity
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                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                              Score 15.4; DB 1; Length 19; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.4; DB 1; Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 T; 0 U;
                                                                                                                                                                                                                                                                                                                                      Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                   0
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RESULT 715
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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                                                                                                                       Biallelic marker; human; high density disequilibrium map; disease; trait; identification; Alzheimer's disease; drug response; drug efficacy;
                                                                                                                                                                                    Human genome biallelic marker primer 220
                                                                                                                                                                                                                              30-JUN-1999
                                                                                                                                                                                                                                                                                                                 AAX52863 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19 BP; 4 A; 10 C; 0 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human anhidrotic ectodermal dysplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             De La Chapelle A, Srivastava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-JUL-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human; anhidrotic ectodermal dysplasia; X chromosome; genetic linkage; translocation; CpG island; foetal development; hair; sweat gland; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUN-1998 (first entry)
                                                                                                    drug toxicity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Col 7; 37pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hair growth.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UNIW ) UNIV WASHINGTON.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-DEC-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US5700926-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            tooth; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo gapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                              339 TTTCCTACCACTCCCCC 355
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16;
                                                                                                                                                                                                                                                                                                                                                                                                                       N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   #5 for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                       TTTCCTACCACTCCACC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STS locus DXS339.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CpG island; foetal development; hair; sweat gland; ss; PCR; amplification; sequence tagged site; STS.
                                                                                                    primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93US-00052997
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                                                                                                                                                                                                                                                                                                                 DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.4;
Pred. No. 8.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4;
. 8.7e+02;
1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Schlessinger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            useful for research into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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WO9904038-A2

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RESULT 716
AAA86302/c
ID AAA863
XX AAA863
XX O4-DEC
XX PCBA H
XX Ribozy
XX Ribozy
XX Mammal
XX MO2000
XX MO2000
XX O6-DEC
XX O4-DEC
XX O4-DEC
XX O4-DEC
XX O4-DEC
XX O1MU-
PA (IMMU-
XX O1MU-
XX
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          constructing a high density equilibrium map of the human genome. The method involves (a) obtaining a nucleic acid library comprising genomic DNA fragments comprising the full genome or a portion (b) determining the order of genomic DNA fragments in the genome, (c) determining the sequence of selected regions of the genomic DNA fragments and (d) identifying nucleotides in the genomic DNA fragments which vary between individuals, thereby defining a set of biallelic markers. The methods can be used for identifying traits such as disease (e.g. Alzheimer's disease), drug response, drug efficacy and drug toxicity. They can be used for selecting an individual for inclusion in a clinical trial. The method is used to map the position of genes in a genome (preferably the human genome). The sequences described in AAX52633-X52832 and AAX52844-X52868 represent primers used in the method of the invention
                                   Tritz R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ribozyme; hairpin; hammerhead; gene therapy; vasotropic; restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA86302;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA86302 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 137; Page 286; 288pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Production of biallelic markers - by obtaining a genomic DNA library, determining the order and sequence of DNA fragments and identifying nucleotides which vary between individuals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18-JUL-1997;
21-APR-1998;
                                                                                                                                                          04-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                  WO200032765-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCBA HH ribozyme binding site #34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                04-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for obtaining a set of biallelic markers represented in AAX52533-X52632 and AAX52833-X52843 for use in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-132278/11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-JUL-1998;
                                                                                                (IMMU-) IMMUSOL INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            910 CAGGGCTCAGAGAGAAG 926
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CAGGGCTCAGAGAGGAG 17
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                                   Welch
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 6 A; 4 C; 8 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97EP-00401740.
98US-0082614P.
                                                                                                                                                          98US-0110954P
                                   Ę,
                                                                                                                                                                                                                       99WO-US028772.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.3%;
                                   Barber JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tchoumakov I;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4; DB 1;
Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                Robbins JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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RESULT 717
AAH61464/c
ID AAH614
XX AAH614
XX AAH614
XX AAH614
XX Human;
KW PCOGIF
KW PTOlif
KW CYTOKI
KW AATISE
CO SYNTHE
XX GOS HOMO
XX Treat
PT That
XX Examp
XX Examp
XX Examp
XX Examp
XX Examp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  문
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; ribozyme therapy; hairpin ribozyme; hammerhead ribozyme; recognition site; target; ribozyme binding site; eye disease; vulnerary; proliferative disease; skin disease; psoriasis; diabetic retinopathy; cytokine; inflammation; cell-cycle dependent kinase; cyclin; MMP; matrix metalloproteinase; growth factor; reductase; scarring; cytostatic; antipsoriatic; dermatological; antiseborrheic; antidiabetic; virucide; antististkling; ophthalmological; keratolytic; gene therapy; viral wart; atopic dermatitis; actinic keratosis; squamous cell carcinoma;
                                                                              Treating proliferative skin or eye diseases that cleave RNA encoding cytokines involved metalloproteinases, growth factors and cell-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a hairpin or hammerhead ribozyme, designed to cleave RNA encoding a cyclin or cell-cycle dependent kinase other than cell-cycle dependent kinases CDK1, PCKN and Cyclin B1. Representative examples of ribozyme recognition sites are given in AAA82415 to AAA86787. The ribozyme of the invention is useful for inhibiting restenosis by introduction of the ribozyme into cells. The ribozyme is resistant to endonuclease activity and hence is efficient
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAH61464;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 19 BP; 5 A; 6 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 105; 109pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New hairpin and hammerhead ribozyme for inhibiting restenosis, cleave RNA encoding a cyclin or cell-cycle dependent kinase other than CDKI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-412314/35
                                                                                                                                                                  WPI; 2001-300427/31.
                                                                                                                                                                                                        Robbins JM,
                                                                                                                                                                                                                                                                                        26-OCT-1999;
                                                                                                                                                                                                                                                                                                                              26-OCT-2000; 2000WO-US029500.
                                                                                                                                                                                                                                                                                                                                                                      03-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                              WO200130362-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCNA HH ribozyme binding site SEQ ID NO:3888.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAH61464 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      restenosis treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCNA and Cyclin B1.
                                                                                                                                                                                                                                                (IMMU-) IMMUSOL INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2328 CACCTTCTTGAAGATGG 2344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cell carcinoma; seborrheic wart; vitreoretinopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CACCTTCTTGAGGATGG
                                                                                                                                                                                                          Tritz R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                        99US-0161532P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            relates to a hairpin or hammerhead ribozyme,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4; DB 1;
Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4; L.
. 8.7e+02;
1;
                                                                                and scarring, using ribo
in inflammation, matrix
-cycle dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 19;
                                                                                                                            using ribozymes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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Example 1; Page 354; 408pp; English.

describes a method

for treating a proliferative

WPI;

2003-679850/64.

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RESULT 718
ADG3560
ID 35660
ID 35660
XX ADG356
XX ADG356
XX ADG356
XX ENA in Well V si
XX Short
XX Synthe
OS Synthe
OS Synthe
OS Human
XX WO200:
XX 20-FEI
XX 20-FEI
XX 21-AU
PP 20-FEI
XX 21-AU
PP 21-AU
PR 21-AU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 16; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                 23-JUL-2002;
21-AUG-2002;
29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleic acid molecule (II) comprising a promoter operably linked to a nucleic acid segment encoding (I). (I) can have antipsoriatic, antisickling, dermatological, cytostatic, antiseborrheic, antidiabetic, antisickling, ophthalmological, vulnerary, keratolytic and virucide activities, and cleaves RNA encoding cytokine involved in inflammation. (I) can be used in gene therapy. (I) and (II) are useful for treating proliferative skin diseases such as psoriasis, atopic dermatitis, actinic keratosis, squamous or basal cell carcinoma and viral or seborrheic wart. They can also be used for treating proliferative eye diseases such as diabetic retinopathy, vitreoretinopathy, sickle cell retinopathy, retinopathy of prematurity and retinal detachment, and for treating and preventing scarring such as keloid, adhesion and hypertrophic or hypertrophic burn scar. AAH57577 to AAH62099 represent sequences used in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    skin or eye disease and scarring. The method involves administering ribozyme (I) which cleaves RNA encoding a cytokine involved in inflammation, matrix metalloproteinase (MMP), cyclin, cell-cycle dependent kinase, growth factor or a reductase, or administering a nucleic acid molecule (II) comprising a promoter creating in the comprision of the comprisi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-FEB-2002;
11-MAR-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         short hairpin RNĀ; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; HIV; anti-HIV; HIV replication inhibition; HIV infection; 8s; target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RNA interference; short interfering nucleic acid; siNA; micro-RNA; miRNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HIV sinh oligonucleotide SEQ ID NO:503.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-APR-2002;
29-MAY-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-FEB-2003;
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                                                  Мсвwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2328 CACCTTCTTGAAGATGG 2344
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunodeficiency virus 1.
                                                                                                                                                        SIRNA THERAPEUTICS INC MCSWIGGEN J.
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2002US-00157580
2002US-0386782P
2002US-0398036P
2002US-00225023
2002US-0406784P
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                                                                                                                                                                                                                                                                                                  2002US-0408378P.
2002US-0409293P.
2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2002US-0363124P.
                                                  Beigelman
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                                                        Macejak
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Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              to a
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RESULT 719
ADG36398/c
ID ADG363
XX ADG363
XX ADG363
XX ADG363
XX ADG363
XX ADG363
XX RNA in
KW RNA in
KW Short
KW Short
KW Short
KW Short
KW HIV re
XX PARIM
XX 20-PEI
PR 20-PEI
PR 20-API
PR 29-ADI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to short interfering nucleic acids (siNA) which CC downregulate expression of HIV gene by RNA interference. The siNAs may or CC may not comprise ribonuclectides and may be double or single stranded. CC They further comprise sense and antisense regions, or alternatively are CC assembled from a sense oligonuclectide and an antisense oligonuclectide. Specifically, the siNAs include short interfering RNA (siRNA), double-CC stranded RNA, micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs CC can be unmodified or chemically modified, can contain CC deoxyribonuclectides, and can be chemically synthesised, expressed from a CC vector or enzymatically synthesised. The invention also relates to kits CC for the in vitro or in vivo delivery of siNA; conjugates and/or complexes CC fiNA, and vectors that express siNA. The siNAs are used to modulate CC expression of the HIV gene in cells, tissue explants or organisms (e.g., CC by ex vivo gene therapy), or in grafts and transplants for the treatment CC used for inhibiting replication of HIV and may be used to treat HIV CC infection. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, CC pharmacogenomics, studying gene function, and gene mapping (e.g., of CC single nucleotide polymorphisms). The present sequence represents the CC upper strand of HIV-targeted double-stranded siNA, which is identical to the HIV transcript target sequence.
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Best Local S
Matches 16
20-FEB-2002;
11-MAR-2002;
12-APR-2002;
22-APR-2002;
29-MAY-2002;
06-JUN-2002;
                                                                                                                                                                                                                                                                                                                                     RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siNNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification; pharmacogenomics; gene function analysis; gene mapping; HIV; anti-HIV replication inhibition; HIV infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 19
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                                                                                                                                     20-FEB-2003; 2003WO-US005190
                                                                                                                                                                                    28-AUG-2003
                                                                                                                                                                                                                            WO2003070193-A2
                                                                                                                                                                                                                                                                        Human
                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HIV sinh oligonucleotide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26-FEB-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADG36398 standard; RNA; 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1261 AGCCTACAGCCCCACCA 1277
                                                                                                                                                                                                                                                                        immunodeficiency virus
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                       ; 2002US-0358580P.
; 2002US-0363124P.
; 2002US-0374722P.
; 2002US-00157580.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 6 A; 9 C; 4 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.3%;
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Pred. No. 8.7e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                            analysis; gene mapping; HIV; anti-HIV; infection; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NO:1241.
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RESULT 720
ADJ66971/c
ID ADJ669
XX ADJ669
XX ADJ669
XX ADJ669
XX ADJ669
XX RET ol
XX RET ol
XX RET ge
KW RET ge
KW multip
XX Uniden
XX JP2003
XX JP2003
PD 30-MAY
XX
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Best Local S
Matches 16
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21-AUG-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              deoxyribonucleotides, and can be chemically synthesised, expressed from a vector or enzymatically synthesised. The invention also relates to kits for the in vitro or in vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the HIV gene in cells, tissue explants or organisms (e.g., by ex vivo gene therapy), or in grafts and transplants for the treatment of a variety of conditions. HIV siNAs have anti-HIV activity. They may be used for inhibiting replication of HIV and may be used to treat HIV infection. The siNAs are also useful for drug screening, diagnosis, therapeutic target identification and validation, genetic engineering, pharmacogenomics, studying gene function, and gene mapping (e.g., of single nucleotide polymorphisms). The present sequence represents the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              may not comprise ribonucleotides and may be double or single stranded. They further comprise sense and antisense regions, or alternatively are assembled from a sense oligonucleotide and an antisense oligonucleotide. Specifically, the sinas include short interfering RNA (siRNA), double-stranded RNA, micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be unmodified or chemically modified, can contain
                                                                                                                                                                                                                                                                                                               ADJ66971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19 BP; 0 A; 4 C; 9 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to short interfering nucleic acids (siNA) which downregulate expression of HIV gene by RNA interference. The siNAs may
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; SEQ ID NO 1241; 170pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pyrimidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New double-stranded short interfering nucleic acid, useful for treating infection with human immune deficiency virus, comprises sugar-modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     15-JAN-2003;
                                                          JP2003156492-A
                                                                                           Unidentified
                                                                                                                               multiple internal
                                                                                                                                                                                                  RET oligonucleotide microchip-related oligonucleotide SeqID74.
                                                                                                                                                                RBT oligonucleotide microchip; mis-sense mutation; mutation hot spot;
                                                                                                                                                gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                       1261 AGCCTACAGCCCCACCA 1277
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
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                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                              immobilised; automatic microarrayer; hereditary cancer;
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2002US-0408378P.
2002US-0409293P.
2002US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 of an HIV-targeted double-stranded siNA.
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2002US-00225023.
                                                                                                                                                                                                                                        (first
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                                                                                                                               secretion
                                                                                                                                                                                                                                                                                                           DNA;
                                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
                                                                                                                             adenomatosis type 2 syndrome;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15.4; DB 1;
No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
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RESULT 721
ADL69619
ID ADL696
XX ADL696
AC ADL696
XX ZO-MAY
XX ZO-MAY
XX RNA iI
KW RNA iI
KW Short
KW Short
KW PARIM
KW PO-0111
XX ZO-PEI
PR 11-FEI
PR 11-SEI
PR 11-SE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                          20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
09-SEP-2002; 2002US-0409393P.
11-SEP-2002; 2002US-0409785P.
11-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention relates to a novel RET oligonucleotide microchip which contains an oligonucleotide that can identify mis-sense mutation of a mutation hot spot in a RET gene, immobilised to the surface of a solid substrate by automatic microarrayer. The RET oligonucleotide microchip is useful for identifying mis-sense mutations in a RET gene, and thus for diagnosing hereditary cancers, where the RET gene is a causative gene of multiple internal secretion adenomatosis type 2 syndromes. The RET oligonucleotide chip enables diagnosis of hereditary cancers accurately and economically. The present sequence is that of an oligonucleotide which is related to the RET oligonucleotide microchip of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RET oligonucleotide microchip contains oligonucleotide that can identify missense mutation of mutation hot spot in a RET gene, immobilized to surface of solid substrate by automatic microarrayer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; short hairpin RNA; shRNA, expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 19 BP; 5 A; 6 C; 5 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 1; SEQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19-NOV-2001; 2001KR-00071774.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19-APR-2002; 2002JP-00117713.
                                                                                                                                                                                                                                                                                                                                                                                18-FEB-2003; 2003WO-US004738.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WO2003070896-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               proliferating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pharmacogenomics; gene function analysis; gene mapping; cancer;
restenosis; cytostatic; vasotropic; human;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human PCNA siNA lower strand,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADL69619;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADL69619 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1771
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2003-692502/66.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18
                                      RIBOZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAT CANCER CENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TGGGTCTTCCAGGAGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TGGGTCTTGCAGGAGCC 1787
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ce11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ID NO 74; 25pp; Japanese.
                                      PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  nuclear antigen; PCNA; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          дB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.4; DB 1;
Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO:78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          micro-RNA; miRNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Activity to the first

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RESULT 722
ADL69550/c
ID ADL695
XX
AC ADL695
XX
DT 20-MAY
XX
DT 20-MAY
XX
Human
XX
RNA in
KW RNA in
KW Short
KW Short
KW Short
KW Short
KW PARMA
W FESTEN
KW PARMA
KW POSO11
XX
V PESTEN
XX
PO 28-AUC
XX
PD 28-AUC
XX
PD 28-AUC
XX
PD 28-AUC
XX
PD 28-AUC
XX
PF 18-FEST
XX
PR 20-FEST
PR 06-JUN
PR 06-JUN
PR 06-JUN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to short interfering nucleic acids (sinh) which CC downregulate expression of the human proliferating cell nuclear antigen CC (PCNA) gene by RNA interference. The sinhs may or may not comprise CC ribonucleotides and may be double or single stranded. They further CC as sense oligonucleotide and an antisense regions, or alternatively are assembled from CC allows and controlled and an antisense oligonucleotide. Specifically, CC micro-RNA (miRNA) and short interfering RNA (siRNA), double-stranded RNA, CC micro-RNA (miRNA) and short hairpin RNA (shRNA). The siNAs can be CC unmodified or chemically modified, can contain deoxyribonucleotides, and CC synthesised. The invention also relates to kits for the in vitro or in CC vivo delivery of siNA, ronjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the PCNA Gene in cells, tissue explants or organisms (e.g., by ex vivo gene CC conditions or They may be used for treating cancer and restenceis. The CC siNAs are also useful for drug screening, diagnosis, therapeutic target CC didnitication and validation, genetic engineering, pharmacogenomics, CC studying gene function, and gene mapping (e.g., of single nucleotide CC polymorphisms). The present sequence represents the lower strand of a human PCNA-targeted double-stranded siNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 11
  20-FEB-2002; 2002US-0358580P.
11-MAR-2002; 2002US-0363124P.
06-JUN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
                                                                                                                                                                                                                                                                                                                                        RNA interference; short interfering nucleic acid; siNA; short interfering RNA; siRNA; double-stranded RNA; micro-RNA; miRNA; short hairpin RNA; shRNA; expression modulation; gene therapy; drug screening; diagnosis; therapeutic target identification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer and restenosis, downregulates expression of the proliferating cell nuclear antigen gene.
                                                                                                                                                                                                                                                                               pharmacogenomics; gene function analysis; gene mapping; cancer;
restenosis; cytostatic; vasotropic; human;
proliferating cell nuclear antigen; PCNA; target sequence; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human PCNA transcript target sequence/siNA upper strand,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 19 BP; 4 A; 4 C; 6 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                               WO2003070896-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20-MAY-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADL69550;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADL69550 standard; RNA; 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-712615/67.
                                                                                                          18-FEB-2003; 2003WO-US004738
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Мсвwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2328 CACCTTCTTGAAGATGG 2344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            μ.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CACCUUCUUGAGGAUGG 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 78; 134pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chowrira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.4; DB 1; Length 19; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO:9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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ARESULT 723
ADN34160/c
ID ADN341
XX ADN341
XX ADN341
XX OPPER
AC OL-JUL
XX Upper
XX Short
KW Short
KW Geneti
KW Geneti
KW Single
XX WO2003
XX WO2003
XX WO2003
XX O4-SEE
PF 06-FEE

short interfering nucleic acid; siNA; cyclin; Cytostatic; Vasotropic; cancer; cell-proliferation disorder; restenosis; drug screening; genetic engineering; pharmacogenomics; gene mapping; single nucleotide polymorphisms; ss.

06-FEB-2003; 2003WO-US003662.

04-SEP-2003 WO2003072705-A2 Homo sapiens Upper strand of cyclin D1 targeted double stranded siNA #180.

01-JUL-2004 (first entry)

ADN34160 standard; RNA; 19

ВP

밁 S

19

CACCITCITGAGGATGG 3

Matches Query Match Best Local

16;

Conservative

0; Mismatches Score 15.4; Pred. No. 8

Similarity

0.3%;

4; DB 1; Length 19; 8.7e+02;

0

0

2328 CACCTTCTTGAAGATGG 2344

Sequence 19 BP; 5 A; 6 C; 4 G; 0 T; 4 U; 0 Other;

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The invention relates to short interfering nucleic acids (siNA) which CC downregulate expression of the human proliferating cell nuclear antigen CPCNA) gene by RNA interference. The siNAs may or may not comprise comprise sense and antisense regions, or alternatively are assembled from CC emprise sense and antisense regions, or alternatively are assembled from CC as bense oligonucleotide short interfering RNA (siRNA), double-stranded RNA, CC micro-RNA (miRNA) and short hairpin RNA (siRNA). The siNAs can be CC unmodified or chemically modified, can contain deoxyribonucleotides, and CC synthesised. The invention also relates to kits for the in vitro or in CC vivo delivery of siNA; conjugates and/or complexes of siNA; and vectors that express siNA. The siNAs are used to modulate expression of the PCNA Gene in cells, tissue explants or organisms (e.g., by ex vivo gene CC therapy), or in grafts and transplants for the treatment of a variety of conditions. They may be used for trasting cancer and restenosis. The CC siNAs are also useful for drug screening, diagnosis, therapeutic target (conditions of function, and gene mapping (e.g., of single nucleotide CC studying gene function, and gene mapping (e.g., of single nucleotide CC polymorphisms). The present sequence represents the upper strand of a transporter tracer sequence sequence represents the permander of a transporter tracer sequence sequence represents the permander of a transporter tracer sequence represents the permander of a transporter tracer sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            05-8EP-2002; 2002US-0408378P.
09-8EP-2002; 2002US-0409293P.
11-8EP-2002; 2002US-0409785P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             proliferating cell nuclear antigen gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New short interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer and restenosis, downregulates expression of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-712615/67.
                                              transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Мсвwiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (RIBO-) RIBOZYME PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 9; 134pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chowrira
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Β,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Beigelman
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RESULT 724
ADN3439
ID ADN343
XX ADN343
XX DI-JUL
XX Short
KW Short
KW Geneti
KW Single
XX SHOMO &
X
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Best Local S
Matches 16
20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
29-AUG-2002;
05-SEP-2002;
09-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-SEP-2002;
09-SEP-2002;
17-SEP-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-FEB-2002;
11-MAR-2002;
06-JUN-2002;
29-AUG-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to a short interfering nucleic acid (siNA) that down regulates expression of at least one cyclin gene by RNA interference. siNA are used to modulate expression of cyclin genes, in cells, tissue explants or organisms, e.g. for treating a wide range of cancers and other cell-proliferation disorders such as restenosis, but also for drug screening, diagnosis, target identification and validation; genetic engineering, pharmacogenomics, studying gene function and gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence represents the upper strand of cyclin D1 targeted double stranded siNA that is the content of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     short interfering nucleic acid; siNA; cyclin; Cytostatic; Vasotropic;
cancer; cell-proliferation disorder; restenosis; drug screening;
genetic engineering; pharmacogenomics; gene mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADN34399;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADN34399 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 3; SEQ ID NO 180; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diagnosis of cancer
                                                                                                                                                                                                                                                   06-FEB-2003;
                                                                                                                                                                                                                                                                                                                                                                                        WO2003072705-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         single nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lower strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JAN-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   5393 ААААААТАСААААА 5409
                                                                                                                                                                                                                                                                                                                                                                                                                                                        sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2003-689983/65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            is identical to the cyclin D1 transcript target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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; 2002US-0363124P.
; 2002US-0386782P.
; 2002US-0406784P.
; 2002US-0408378P.
; 2002US-0409293P.
; 2002US-0411275P.
; 2003US-0440129P.
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ilarity 94.1%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of cyclin D1 targeted double stranded siNA #180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polymorphisms; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RNA;
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Pred. No. 8.7e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other;
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RESULT 725
AAQ95360
ID AAQ953
XX AAQ
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Best Local S
Matches 15
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                                                                                                  Kit for automated genotyping contg. pairs of PCR primers - de amplify polymorphic nucleotide repeat sequences, arranged in with a characteristic fluorescence label, useful e.g. in dete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer; polymerase chain reaction; PCR; linkage study; locus; microsatellite marker sequence; automated genotyping; allele; polymorphism; detection; Homo sapiens; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Primer B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   08-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAQ95360 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           one cyclin gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New short interfering nucleic acid, useful e.g. for treatment diagnosis of cancer and restenosis, down regulates expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Thompson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-SEP-2002; 2002US-0411275P.
15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                     WPI; 1995-215278/28.
                                                                                                                                                                                                                                                                          Levitt
                                                                                                                                                                                                                                                                                                                                                                                  03-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                    05-DEC-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9515400-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a short interfering nucleic acid (that down regulates expression of at least one cyclin gene by RNA interference. siNA are used to modulate expression of cyclin genes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 3; SEQ ID NO 419; 144pp; English
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                                                                                                                                                                                                                                                                                                                            (UYJO ) UNIV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (Group 1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP; 14 A; 2 C; 2 G; 0 T; 1 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mcswiggen J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                               JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                  93US-00160837.
                                                                                                                                                                                                                                                                                                                                                                                                                                       94WO-US013945.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set C)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
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Pred. No. 8.7e+02;
1; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                             designed to in sets each letection of
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Claim 3;

Fig 7A-2; 104pp; English

related genetic

rearrangement.

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RESULT 726
ANT27911/c
ID ANT279
XX AAT279
XX AAT279
XX Detect
KW Detect
KW adapto
KW Beneti
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX Hollor
PN W09617
XX O6-JUN
XX CDUPO
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Best Local Similarity
Matches 16; Conserv
Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods. to detect differences. The primers used in the amplification comprise a primer consisting of a perfect cpd. simple sequence repeat (SSR), and an adaptor directed primer, comprising a sequence complementary to an adaptor directed primer, comprising a sequence of a SSR primer, which is segment. The present sequence is an example of a SSR primer, which is flanked at its 5'-end by degenerate nucleotides. The method represents a modified amplified fragment length polymorphism assay, which is partic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The method aims to provide a collection of highly reproducible microsatellite marker sequences (MMS) at approx. 10-50 cM intervals throughout the human genome which can be detectably labelled. The MSS are polymorphic, simple sequence repeats and can be used in automated genotyping, esp. fluorescence-based. The primers correspond to the unique DNA sequence surrounding each marker, and PCR is used to detect each polymorphism. When the MSS show considerable polymorphism (ie. a difference in the number of repeats) between individuals, the markers can be particularly informative. The MSS can be ideal for linkage studies. Kits comprise at least 4 groups, of at least 3 sets, each comprising labelled primers for PCR amplification of the DNA. Group 1 primer pairs are shown in AAQ95327-68. The published size range of the IGP1 allele is 176-196 bp, and the degree of heterozygosity in the population is about
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modified amplified fragment length polymorphism assay -polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite region; genetic trait marking; germplasm comparisons; 5'-anchored; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Morgante M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-NOV-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Page 76; 173pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PONT DE NEMOURS &
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vogel
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Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                       Detecting polymorphisms between 2 nucleic acid samples, esp. in microsatellite regions, comprises digesting the nucleic acid to generate fragments, ligating adaptor segments to their ends, amplifying them using primer directed amplification and comparing the prods to detect differences. The primers used in the amplification comprise a primer consisting of a perfect pod, simple sequence repeat (SSR), and an adaptor directed primer, comprising a sequence complementary to an adaptor directed primer, comprising a sequence of a SSR primer, which is flanked at its 5'-end by degenerate nucleotides. The method represents a modified amplified fragment length polymorphism assay, which is partic. useful for genome fingerprinting, i.e. for genetic trait marking and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detection; polymorphism; perfect compound simple sequence repeat; adaptor directed primer; genome; genetic; fingerprinting; amplified fragment length polymorphism assay; microsatellite regigenetic trait marking; germplasm comparisons; 5'-anchored; ss.
                                                                                                                                                                    Sequence
                                                                                                                                                                                                                    germplasm comparisons
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Modified amplified fragment length polymorphism assay polymorphism esp. in micro:satellite regions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1996-277795/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
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                                                          16; Conserv
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                                                                                                                                                                 BP; 8 A; 0 C; 9 G; 0 T; 0 U; 3 Other
                                                             Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                   94.18;
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                                                                                      Score 15.4; DB 1; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.4; DB 1
Pred. No. 8.7e+02
                                                             Mismatches
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                                                                                                                  DB 1;
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                                                                                                            Length 20
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RESULT 729
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AAT6579
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Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  markers. Primers based on these sequences can be used to detect these repeats, especially for use in e.g paternity or maternity testing, human genetic analysis such as linkage analysis of genetic disease, commercial animal or plant breeding or pedigree analysis. Clones containing the repeat sequences were isolated by hybridisation of chromosome-specific phage libraries with a synthetic poly(dC-dA), (dG-dT) probe. Over 100 repeat blocks were isolated. The primers AAT65798-T66047 were used to PCR amplify the inserts from the isolated clones containing the repeat sequences. The primers AAT65798-9 were used to amplify the repeat sequence marker clone Mfd1 (AAT65797). (Updated on 25-MAR-2003 to correct
                                Construction
                                                                                                                                             AAT97382
                                                                                                                                                                                                  AAT97382 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to the isolation of polymorphic repeat sequences having the sequence (dC-dA)n.(dG-dT)n which can be used as genetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detection of polymorphic genetic markers of the form (dC-dA)n(dG-dT)n using novel nucleic acid mols. as primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-APR-1989;
05-SEP-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         linkage analys
hybridisation;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           25-MAR-2003
17-JUN-1997
                                                                                     23-MAR-1998
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                             of a
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                                                                                     (first entry)
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(first entry)
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91US-00754351.
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                          GALV retroviral packaging plasmid using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A; 4 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                    DNA;
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                                                                                                                                                                                                    ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.4; DB 1; Length 20; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                             primer 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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AAT92764/c
ID AAT9277
XX AAT927
XC AAT927
XX D5-PEB
XX Primer
XX PCR pr
KW PCR pr
KW transg
KW Huntin
KW myelom
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                                                                                                                                                                                                                                                                                                                             S
                                                                                                                                                                                                                                                                                                                                                                                                                                          CY This primer is used in the construction of a retroviral packaging plasmid CY which contain genes encoding GALV gag/pol or envelope protein. The CY retroviral packaging plasmid is used for the production of high titres of CY recombinant retrovirus in human cells. The retroviral packaging plasmid CY comprises a retroviral helper DNA sequence derived from a replication-CY incompetent retroviral helper DNA sequence derived from a replication-CY comprises a ecotropic Moloney murine leukaemia virus (MMLV) or gibbon ape CY encodes a ecotropic Moloney murine leukaemia virus (MMLV) or gibbon ape CY cectropic or polytropic envelope protein. The packaging plasmids, are used with a second retroviral vector CY designated KAT plasmids, are used with a second retroviral vector CY cectropic or polytropic envelope protein. The packaging plasmids with CY retroviral supernatants that express, e.g. a hormone, lymphokine, growth CY factor or coagulation factor. The plasmids are useful in construction of stable cell lines that constitutively produce the retroviral proteins CY required in trans for the production of retrovirus particles: gag, pol CY and env. The stable producer cells continue to produce high titre CY ertroviral vector plasmids are constructed with sequences enabling the cyleron all persistence without the need for stable integration of the production of retrovirus particles.
                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 16; Conserv
PCR primer; amplify; human gene; chimeric non-human animal; antibody; transgenic mouse; chromosome fragment; hybridoma production; microcell; Huntington's disease gene; pluripotent cell; interleukin-2 gene; myeloma cell; immunoglobulin; variable region; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  High efficiency retroviral packaging s cells, esp. haematopoietic stem cells,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
Gibbon ape leukemia virus
                                                                                        Primer #1 for immunoglobulin kappa variable region Vkappa1-2
                                                                                                                               05-FEB-1998
                                                                                                                                                                 AAT92764;
                                                                                                                                                                                                     AAT92764 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 14; Page 92; 157pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-165307/15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-AUG-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-AUG-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              retroviral packaging system; human; virion protein; expression vector; PCR primer; recombinant; GALV_f ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            vector plasmid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CELL-) CELL GENESYS INC.
                                                                                                                                                                                                                                                                                                                        4821 CACCAGCCCTTGACCCT 4837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ŧ
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                                                                                                                                                                                                                                                                                             CACCAGCCCATGACCCT 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dull TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 2 A; 2 C; 11 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                               (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-00517488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96WO-US013737.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The Env gene determines the host range
                                                                                                                                                                                                     DNA;
                                                                                                                                                                                                                                                                                                                                                                                  0.3%;
                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zsebo
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                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                    Score 15.4; DB 1
Pred. No. 8.7e+02
                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cooke K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    system
s, T or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Farson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      B '
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  used to transduce human cells with foreign genes.
                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
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                                                                                                                                                                                                                                                                                                                                                                       Indels
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RESULT 731
AAV49811
ID AAV498
XX AAV498
XC AAV498
XC AAV498
XX ACTIVI
KW ACTIVI
KW ACTIVI
KW SPINAI
KW N-meti
KW N-meti
KW N-meti
KW N-meti
KW N-meti
KW Synthi
OS Synthi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC AAT92758-T92817 represent amplification primers for human genes which are CC used in the chimeric non-human animal of the invention. The chimeric non-human animal of the invention. The chimeric non-cC human animal of the invention, preferably a mouse, contains a foreign chromosome(s) or chromosome fragment. The animal is produced by obtaining the hybrid cell by fusion of a cell containing the foreign chromosome with a cell having the ability to form microcells. The microcells are crompared, and fused with cells having differentiative pluripotency to form cells having differentiative pluripotency and containing the foreign chromosome. These cells are then introduced into an embryo, which is then containted and brought to term. The foreign for an antibody. The containted and brought to term. The foreign for an antibody. The containted and brought to term. The foreign for an antibody. The containted and brought to term. The foreign for an antibody. The containted and the Huntington's disease containted and as the interleukin-2 gene, and the Huntington's disease containted and as the interleukin-2 gene, and the Huntington's disease containted and as the interleukin-2 gene, and the Huntington's disease containted and as the interleukin-2 gene, and the Huntington's disease containted and as the interleukin-2 gene, and the Huntington's disease containted and the Huntington's disease containted and containted and fused with the produce hybridoms capable of expressing the foreign capable of expr
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Best Local S
Matches 16
                                                                 Activity dependent neurotrophic factor III: ADNF-III; ADNP; cell deat activity dependent neuroprotective protein; neurone; excito-toxicity; spinal cord; hippocampus; cerebral cortex; cholinergic; beta-amyloid; N-methyl-D-aspartate; Alzheimer's disease; human immunodeficiency vir HIV infection; PCR primer; ss.
                                                                                                                                                                                                                                             04-NOV-1998
    Synthetic.
Homo sapiens.
                                                                                                                                                                                                                                                                                        AAV49811;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 1; Page 21; 142pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric animal containing foreign chromosome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1997-178822/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-AUG-1995;
15-PEB-1996;
                                                                                                                                                                                                   ADNF-III PCR primer #3.
                                                                                                                                                                                                                                                                                                                              AAV49811 standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tomizuka K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              06-MAR-1997.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (KIRI ) KIRIN BEER KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     865 GCAGTGCTAATGCCCTG 881
                                                                                                                                                                                                                                                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gene, e.g. an antibody.
                                                                                                                                                                                                                                                                                                                                                                                                                                          GCACTGCTAATGCCCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to produce the antibody)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                             (first
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96JP-00027940.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; 4 C; 7
                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.3%;
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                                                                                                                                                                                                                                                                                                                                20
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                                                                                                                                                                                                                                                                                                                              ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 4 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 15.4; DB 1;
Pred. No. 8.7e+02;
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0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 20;
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                                                                                                                                                        cell death;
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RESULT 732
AAV52761/c
ID AAV527
XX AAV527
XX AAV527
XX Immuno
XX Immuno
XX Plurip
KW Plurip
KW Constr
XX Synthe
OS Homo s
XX WO9837
XX WO9837
XX O3-SEP
XX O2-MAR
XX A (KIRI
XX XIRI
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV49809-V49811 are PCR primers used in the amplification of a novel activity dependent neurotrophic factor III, ADNF-III (also known as activity dependent neuroprotective protein, ADNF). ADNF III polypeptides can be used to prevent neuronal cell death, of e.g. the spinal cord, hippocampal, cerebral cortical or cholinergic neurones associated with e.g. HIV infection, excito-toxicity induced by N-methyl-D-aspartate stimulation or beta-amyloid peptide in Alzheimer's disease. The polypeptides can also be combined with a carrier to alleviate learning impairment produced by cholingeric blockage in Alzheimer's patients. The nucleic acids are useful in polypeptide production and to detect ADNF III polynucleotide in biological samples, while the antibodies are useful therapeutically and to isolate ADNF III polypeptides
                                             28-FEB-1997;
                                                                        02-MAR-1998;
                                                                                                      03-SEP-1998
                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                    Pluripotent cell; intrinsic gene; chimeric non-human animal; construction; human; antibiotic gene; cancer cell; embryonic
                                                                                                                                                                                                                                                               Immunoglobulin kappa variable PCR primer Vk1-2
                                                                                                                                                                                                                                                                                            27-NOV-1998
                                                                                                                                                                                                                                                                                                                         AAV52761
                                                                                                                                                                                                                                                                                                                                                    AAV52761 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 9 A; 6 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 10; Page 5; 121pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Activity dependent neurotrophic factor III polypeptide - useful therapeutically to prevent neuronal cell death associated with e.g. HIV infection, excito-toxicity or Alzheimer's disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-FEB-1997;
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                  (KIRI ) KIRIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                      3305 АССТССАССАСЛАСЛАС 3321
                                                                                                                                                            sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                            μ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                            (first entry)
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                                              97JP-00062309
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                                                                           098000dr-0M86
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                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.4; DB 1;
Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                     embryonic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0
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learning
                                                                                                                                                                                                                        PCR primer;
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RESULT 733
AAX79722
ID AAX797
XX AAX797
XX AAX797
XX 17-AUG
XX PCR px
XX JP1111
XX Geneti
PT Geneti
PT Geneti
PT Geneti
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CC The present invention describes a method of obtaining pluripotent cells CC containing foreign chromosomes or their fragments (preferably at least CC 670 kb in length, especially more than 1000 kb) by preparing cancerous cells containing the foreign chromosomes or fragments, then fusing these CC with pluripotent cells such as embryonic stem cells, embryonic creproductive cells, embryonic cancer cells or their mutants. Also described are: (1) a method of obtaining hybridoma cells by fusing a cell with a high ability to produce hybridoma cells (such as mouse A9 cells) cowith a cell containing the foreign chromosomes or fragments (such as normal human diploid cells); (2) a method of utilising pluripotent cells to produce chimeric and transgenic non-human animals (especially mammals such as mice) which can express the foreign chromosomes or fragments cells contained to an express the foreign chromosomes or fragments cells contained to an express the foreign chromosomes or fragments cells contained to an express the foreign chromosomes or fragments cells controduced, and (3) chimeric animals, their offspring and tissues and cells derived from the offspring produced by a method as in (2). The control of the contained cells contained the production of chimeric and contained the production of chimeric and contained the study of human type and therefore not antigenic in serve as models for the study of human diseases. AAV52755 to AAV52828 are CC primers used in examples from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 16; Conserv
                                         Genetic diagnosis using human mitochondrial DNA -
                                                                                                                                                                  13-OCT-1997;
                                                                                                                                                                                                       13-OCT-1997;
                                                                                                                                                                                                                                               27-APR-1999
                                                                                                                                                                                                                                                                                      JP11113597-A
                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-AUG-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAX79722;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAX79722 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 5 A; 4 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pluripotent cells containing foreign chrohuman chimeric animals constructed using genes such as human antibiotic genes.
                                                                                   WPI; 1999-320841/27
                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer L5582 for mitochondrial DNA analysis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example
                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer; human; mitochondrial DNA; genetic diagnosis;
                        replacements
                                                                                                                                                                                                                                                                                                                            sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              865 GCAGTGCTAATGCCCTG 881
                                                                                                                                                                                                                                                                                                                                                                                        disease contraction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                          TANAKA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GCACTGCTAATGCCCTG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Page 34; 217pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                          3
                                                                                                                                                                  97JP-00279127
                                                                                                                                                                                                       97JP-00279127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
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                                                                                                                                                                                                                                                                                                                                                                                        88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.4; DB 1; Length 20; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chromosomes or fragments - and non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oshimura M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           them
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and expressing foreign
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                         comprises detecting
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This sequence represents a PCR primer that can be used in the method of the invention. The method is for genetic diagnosis using human mitochondrial DNA where there is at least one base replacement from among the following five replacements: the 3010th base is changed from guanine to adenine; the 4883rd base from cytosine to thymine; the 5178th base from cytosine to adenine; the 8414th base from cytosine to thymine; and the 14688th base from cytosine to thymine. The method can be used for diagnosing the probability of contracting adult diseases. A confirmation of base replacement can give a diagnosis of the level of probability of
                                                 AAX91991-X97517 represent PCR primers used to amplify open reading frames and other nucleic acid sequences from the genome of Chlamydia pneumoniae (see AAX91990). C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAX91584- AAX95879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising
                                                                                                                                                                                                                                                                                                                                     Griffais R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis; sinusitis; purulent otitis media; erythema nodosum; pharyngitis; vaccine; neutralising epitope; PCR primer; ss.
                                                                                                                                                                                                                                                                    Genome sequence of Chlamydia pneumoniae
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04-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chlamydophila pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PCR primer used to amplify an ORF of Chlamydia pneumoniae
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAX93323;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      contraction of adult diseases
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                                                                                                                                                                                                                                                                                                                                                                        (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3070 ACAGCTGAGGACTGCAA 3086
                                                                                                                                                                                                                                     1580; Disclosure;
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nilarity 94.1%;
Conservative
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98US-0107078P.
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                                                                                                                                                                                                                                    1912pp; English.
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Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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Sequence 20

BP; 6 A; 7

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                             cc expression of the gene, and an effective symmetric amount of a protein cc effector of a product of the gene. Also described are: (1) a method for treating a disease responsive to inhibition of a gene in a mammal; (2) a cm thod for inhibiting tumour growth in mammal; (3) an inhibitor of a gene comprising an antisense oligonucleotide which inhibits expression of the comprising an antisense oligonucleotide which inhibits expression of the cand (4) a pharmaceutical composition comprising the inhibitor of (3). The cand (4) a pharmaceutical composition comprising the inhibitor of (3). The cut of the gene in composition are useful as analytical tools for transgenic studies and as therapeutic tools, e.g. as gene therapy tools for human cdiseases including benign and malignant tumours, inflammation or asthma. CC expression or activity of a gene or gene product may be used to treat compatients having, or predisposed to developing, a disease responsive to provide missing gene functions and improve a given condition. CC Furthermore, the methods and compositions are useful as probes of the composition of a gene product in an experimental cell culture or animal system; and to evaluate the effect of inhibiting gene activity or expression. AAA55758 to AAA55942 represent oligonucleotide sequences
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; DNA methyltransferase; DNA MeTase; antisense oligonucleotide; modulation; inhibition; gene expression; combination therapy; p16; histone deacetylase; HDAC; thymidylate synthase; tumour suppressor; methylation; gene therapy; tumour; cytostatic; antiasthmatic; antiinflammatory; inflammation; asthma; ss.
                                                                                                                                                                                                                                                                                                                                                                        a gene in a cell comprising contacting the cell with an effective synergistic amount of an antisense oligonucleotide which inhibits
                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                effectors e.g. 5-aza-cytidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Besterman JM,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                with a synergistic
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                                                                                                                                                                                                                                                                                                                                                                                                          invention describes a method for inhibiting the expression
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              n e.g. DNA methyltransferase, of antisense oligonucleotide dine of gene products, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HD2 antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Siders WM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.4; DB 1;
Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              by treating cells and protein for gene therapy
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The present sequence is the sense PCR primer, that corresponds to 71-90 CC bases of mouse Activity Dependent Neurotrophic Pactor (ADNF) III cDNA, CC used for amplification of cDNA from human neuroblastoma. ADNF III is also CC called an Activity Dependent Neuroprotective Protein (ADNP) and is CC expressed in the astrocytes, brain and also in foetal lung and endocrine CC tissues. ADNF III has homology to ADNF II and hsp60, a heat shock protein CC end pIF1, a DNA repair protein. The ADNF III polypeptides are useful for the treatment of neurological deficiencies and for prevention of neuronal CC cell death associated with 99120, the envelope protein from HIV, N-methyl CC degeneration acid (excito-toxicity); tetrodotoxin (blockage of electrical activity); and beta-amyloid peptide, a substance related to neuronal CC degeneration in Alzheimer's disease. It is useful for the treatment of CC Huntington's disease, AIDS dementia complex, epilepsy, neuropathic pain CC mitochondrial abnormalities, Leber's disease, Wernicke's encephalopathy, allcheimer's disease, homocysteinuria, hyperprolinemia, sulphite oxide CC disease, Tourette's syndrome, oxidative stress induced neuronal death, drug Down's syndrome, developmental retardation and learning impairments, drug
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADNP; Activity Dependent Neuroprotective Protein; neuronal cell death; HIV; Human Immunodeficiency Virus; neurological deficiency; treatment; Alzheimer's disease; beta-amyloid peptide; Huntington's disease; epilepsy; AIDS dementia complex; neuropathic pain syndrome; ALS; amyotrophic lateral sclerosis; Parkinson's disease; Leber's disease; mitochondrial abnormality; Wernicke's encephalopathy; homocysteinuria; hyperprolinemia; sulphite oxide disease; Tourette's syndrome; nootropho Down's syndrome; drug addiction; developmental retardation; antilipemilearning impairment; anticonvulsant; neuroprotective; anti-HIV; ss.
                                                                                                                                                                                                                                                                                                                                                                                                   New nucleic acid encoding an activity dependent neurotrophic factor III (ADNF III) useful in the treatment of neurological deficiencies and for preventing neuronal cell death.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; 866T-AON-90
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sense PCR primer #2, to amplify ADNF III cDNA from human neuroblastoma
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                                                                                                                                                                                                                                                                                                                                                              Claim 10; Page 91; 136pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (USAS )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GOVERNMENT US REPRESENT AS.
UNIV RAMOT APPLIED RES & IND DEV LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.4; DB 1;
Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      syndrome; nootropic;
rdation; antilipemic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PCR primer;
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                                                                                                            Query Match
Best Local Similarity
Matches 16; Conserv
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Best Local Similarity
                                                                                                                                                                                                                                                                                The invention relates to a novel method of producing cells containing a modified foreign chromosome or chromosome fragment. The method comprises:
(a) fusing a microcell comprising the foreign chromosome or chromosome fragment, with a cell having a high efficiency for homologous recombination; (b) marking the desired site of insertion of the foreign chromosome using a targeting vector; and (c) inducing deletion or translocation at the marked site. Transgenic animals produced by the method are useful to provide disease models and knockout animals, and in the production of human proteins, particularly human antibodies. This sequence is used in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 9 A; 6 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 1; Page 55; 316pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Producing a cell containing modified foreign chromosomes, useful for the generation of transgenic animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       generation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Primer 1 for human immunoglobulin kappa variable region gene Vk1-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-JUL-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (KIRI ) KIRIN BEER KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       primer; human; ss.
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                                    865 GCAGTGCTAATGCCCTG 881
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20 GCACTGCTAATGCCCTG 4
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                                                                                                                                                                                                                               BP; 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98JP-00236169
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                                                                                                                                                                                                                               A; 4 C; 7 G; 4 T; 0 U; 0 Other;
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                                                                                                                                           0.3%;
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                                                                                                                                           Score 15.4; DB 1; Length 20; Pred. No. 8.7e+02;
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Pred. No. 8.
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RESULT 738 AAA80308/c

Sequence 20

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CC family of transcription factors, which have been implicated in the genes known to be important in the aetiology of asthma. Both ASTHII and CR ASTHIJ mRNAs are alternatively spliced. Alternative splicing of CC ASTHIJ mRNAs are alternatively spliced. Alternative splicing of CC transcripts has no effect on the open reading frame of ASTHIJ, as the CC exons involved are all 5' to the start codon in exon b. In contrast, CC alternative splicing of ASTHII transcripts results in 3 different ASTHII CC alternative splicing of ASTHII transcripts results in 3 different ASTHII CC alternative splicing of ASTHII transcripts results in 3 different ASTHII CC alternative splicing of ASTHII transcripts results in 3 different ASTHII CC predisposition to asthma, as probes for identifying ASTHI related genes, CC for identifying expression of the gene in a biological specimen, and for CC generating genetically modified non-human animals or site specific gene compositions that mimic or modulate activity or expression of ASTHII consociates in drug screening for CC compositions that mimic or modulate activity or expression of ASTHII CC compositions that mimic or modulate activity or expression of ASTHII CC compositions are useful at the introduce proteins; and an animal so received as a consociate specific and therapeutic purposes. The intact ASTHII or ASTHII consociates are useful in the identification of gene activity in vivo for CC development of asthma, and for modulation of gene activity in vivo for proteins or active fragments thereof may be used to modulate or reduce transcript related the consociate the activity. Sequences AABOC60-ABOC61 and AAABOC64-ABOC61 and AAABOC64-ABO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ASTH1 protein, for function of Figure 1 predisposition to the predisposition to the predisposition to the predisposition to the protein for the protein for the protein for the protein for the protein, for the protein for the protei
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          hyperreactivity. The ASTH1I and ASTH1J genes are oriented in opposite directions with the ASTH1 locus, and have similar patterns of expression and common sequence motifs. They are both expressed in trachea, lung and several other tissues. ASTH1I and ASTH1J are novel members of the ets
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to the ASTH1 locus on the short arm of human chromosome (11p). This locus comprises the ASTH11 and ASTH1J genes, where associated with a genetic predisposition to asthma and bronchial hyperreactivity. The ASTH1I and ASTH1J genes are oriented in opposite.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acids other than naturally occurring chromosomes ASTH1 protein, for e.g. screening compositions that modulate or function of ASTH1 proteins or as diagnostics for genetic
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Brooks-Wilson
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01-JUL-1997;
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                                                                    polymorphic
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97US-0051432P
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larey AH;
                                                                        sites within
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                                                                        the ASTH1J
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RESULT 739
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XX AAT19e
XX AAT19e
XX AAT19e
XX Ecll F
XX Gell Gell
XX Gell
XX Gell Gell
XX Gell Gell
XX Gel
RESULT 740
AAF54889
ID AAF548
XX
AC AAF548
XX
AC AAF548
XX
DT 15-MAY
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Best Local S
Matches 16
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The sequences given in AAH43115-21 are oligonucleotides which are antisense to the histone deacetylase gene, HDAC-2. These oligonucleotides may be used in combination with an inhibitor of histone deacetylase enzyme function, to given an improved inhibitory effect, thereby reducing the amount of inhibitor required to obtain a given inhibitory effect. Compounds containing these oligonucleotides may be used to treat cell proliferation conditions such as cancer, restenosis or psoriasis. They can also be used to treat protozoal and fungal infections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New inhibitors of histone deacetylase e.g. N-hydroxy-5-(4-(benzenesulfonylamino)-phenyl)-4-yn-2-pentanamide for treating cancer, restenosis or fungal infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           31-MAY-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cell proliferation; ca fungal infections; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antisense; histone deacetylase; HDAC-1; cell proliferation; cancer; restenosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-SEP-2001
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                             15-MAY-2001
                                                                             AAF54889;
                                                                                                                                AAF54889 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 40; 147pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Delorme D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-NOV-2000; 2000WO-IB001881.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAH43117;
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                                                                                                                                                                                                                                                                                                 2640 CCTGCAGCTGCTGC 2656
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4891 ACAAGTTGCCATCTGGTTT 4909
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                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99US-0167035P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; 9 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                DNA;
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                             entry)
                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
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Pred. No. 8.7e+02;
1; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                          Score 15.4; DB 1; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Thibault C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Abou-Khalil
                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20
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CC PCR primers AAF54889-90 were used to amplify human activity dependent concurrophic factor (ADNP) III cDNA. The primers are based on murine ADNP CIII. ADNP III is involved in prevention of neuronal cell death, and CC foetal alcohol syndrome. The specification describes a method for comprises contacting the growth of a pathologically proliferating cell. The method comprises contacting the cell with an ADNP III antisense oligonucleotide. CC pathologically proliferating or malignant cells, e.g. breast cancer, curroblastoma, ovarian cancer, endometrial cancer, prostate cancer, colladder cancer, lung cancer, oseophageal cancer, prostate cancer, colladder cancer, colon cancer, cosophageal cancer, pancreatic cancer or CC leukaemia. The method may also be used to inhibit and/or detect the CC growth of benign proliferating cells, e.g. restenotic plaques in vascular smooth muscle (e.g. atheroselerosis or restenotic plaques in vascular cancer method may be used to inhibit the growth of cells, retinal hemangioblastomas or psoriatic cells. The cell method may be used to inhibit the growth of cancer cells ex vivo or in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gozes I,
Bassan M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Inhibiting the growth of malignant or benign proliferating cells (e.g. breast cancer or psoriatic cells, respectively), by contacting the cells with an activity dependent neurotrophic factor III antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   neuronal cell death; foetal alcohol syndrome; proliferating cell; restenotic plaque; atherosclerosis; restenosis; retinal hemangioblastoma; benign prostatic hyperplasia cell; PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 29; 57pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-159867/16.
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(USSH )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Activity dependent neurotrophic factor III; ADNF III; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR primer for activity dependent neurotrophic factor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              UNIV RAMOT APPLIED RES & IND DEV LTD US DEPT HEALTH & HUMAN SERVICES.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Zamostiano R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gelber
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Sequence 20 BP; 9 A; 6 C; 2 G; 3 T; 0 U; 0 Other;

Ś Query Match Best Local S Matches 16 3305 l Similarity 16; Conserv ACCTGCAGCAGAACAAC 3321 0.3%; nilarity 94.1%; Conservative 0; Score 15.4; DB 1; Pred. No. 8.7e+02; Mismatches Length Indels 20; 0 Gaps

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ACCTGCAGCAAAACAAC 17

RESULT 741
AAC89537
ID AAC895
XX
XX
AC AAC895
XX
DT 08-MAR
XX
DB Human
XX
KW Histon
KW Histon Human HDAC-2 PCR primer SEQ ID NO: 7. AAC89537 AAC89537 standard; DNA; (first entry) 20 ВP

Histone deacetylase; HDAC-1; HDAC-2; HDAC-3; HDAC-4; HDAC-5; HDAC-C; HDAC-D; cell cycle; tumourigenesis; cancer; inhibitor; antisense;

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RESULT 742
AAC89546
ID AAC895
XX AAC895
XX OB-MAR
AC OB-MAR
XX Histon
KW HDAC-D
KW Gene t
XX Gene t
XX Gene C
XX Gene
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention provides inhibitors of histone deacetylase enzymes such as HDAC-1, HDAC-2, HDAC-3, HDAC-4, HDAC-5, HDAC-C and HDAC-D. These inhibitors may be antisense strands or they may be compounds identified by contacting the enzyme with the compound and measuring the resulting enzyme activity. These inhibitors are useful for treating cancers and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Histone deacetylase; HDAC-1; HDAC-2; HDAC-3; HDAC-4; HDAC-5; HDAC-C;
HDAC-D; cell cycle; tumourigenesis; cancer; inhibitor; antisense;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antisense oligonucleotide that inhibits expression of a histone deacetylase, useful for treating and/or alleviating the symptoms neoplasia, or for inhibiting neoplastic cell growth in an animal
                                                  WPI; 2001-016407/02
                                                                                                                                                                                                                                                            03-MAY-2000; 2000WO-IB001252
                                                                                                                                                                                                                                                                                                              30-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                WO200071703-A2
                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human HDAC-2 antisense sequence SEQ ID NO: 16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAC89546;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 0 A; 9 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          enzyme activity. These inhibitors are userus to tracery continuity which histone deacetylase is involved in a neoplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 12; 125pp; English
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                                                                                                       Macleod
                                                                                                                                                         (METH-) METHYLGENE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (METH-) METHYLGENE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2001-016407/02.
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     oligonucleotide that inhibits
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                  Li Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Li Z,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                       Besterman
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Pred. No. 8
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  expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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     histone
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RESULT 743
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Best Local S
Matches 15
                                         The invention is directed to BAP28 polypeptides, BAP28 polynucleotide sequences and regulatory region located at the 3' and 5' ends of the BAP28 coding region. The BAP28 polypeptides can be expressed by standard recombinant methodology. BAP28 polypucleotides and polypeptides have been found to be over expressed in prostate tumour cells, therefore levels of BAP28 expression and/or activity may be assayed (e.g. by polymerase chain reaction (PCR)) to diagnose patient suffering from or susceptible to prostate cancer. Antibodies specific for the BAP28 polypeptides are useful as diagnostic reagents. Biallelic markers of the BAP28 gene are useful in genetic analysis. Sequences AAP83934-963 represent primers for the BAP28 gene and PCTA-1 gene (the coding strand of PCTA-1 gene is on the BAP28 gene and PCTA-1 gene (the coding strand of PCTA-1 gene is on
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention provides inhibitors of histone deacetylase enzymes such as HDAC-1, HDAC-2, HDAC-3, HDAC-4, HDAC-5, HDAC-C and HDAC-D. These inhibitors may be antisense strands or they may be compounds identified by contacting the enzyme with the compound and measuring the resulting enzyme activity. These inhibitors are useful for treating cancers and for identifying which histone deacetylase is involved in a neoplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       deacetylase, useful for treating and/or alleviating the symptoms neoplasia, or for inhibiting neoplastic cell growth in an animal
                                                                                                                                                                                                                                                                              New BAP28 polynucleotides and polypeptides overexpressed in prostate cancer cells for diagnosing prostate tumors, e.g. by hybridization or polymerase chain reaction assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                 Barry C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BAP28 gene fragment amplifying primer BAP28polyTcourt
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAF83959;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 24; 125pp; English.
                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-367032/38
                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-JUN-1999; 99US-0141323P.
18-JAN-2000; 2000US-0176880P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-JAN-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BAP28; prostate; tumour; cancer; diagnostic; genetic analysis; PCTA-1;
                                                                                                                                                                                                                                               Example; Page 347; 349pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                     (GEST )
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP; 0 A; 9 C; 5 G; 4 T; 2 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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Pred. No. 8.7e+02;
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opposite of the coding

strand of BAP28)

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0 Other

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RESULT 744
AAD12482/c
AAD12482;
XX
AAD12482;
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AAD12482;
XX
CAAD12482;
XX
DE Mouse caspase 8 mRNA antisense (
XX Caspase 8; infection; inflammati
XW Gene therapy; antisense; mouse;
XX
Mus musculus.
OS Synthetic.
XX
CASPASE 1..20
PT modified_base 1..20
PT modified_base 1..20
PT modified_base 1..5
PT modified_base 2/tag= a
PT modified_base 1..5
PT modified_base 2/tag= a
PT modified_base 3/tag= d
PT modified_base 5/tag= g
PT modified_base 5/tag= g
PT modified_base 11
PT modified_base 12
PT
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Best Local Similarity
Matches 16; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           infection; inflammation; tumour; research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.3%;
ilarity 94.1%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antisense; mouse; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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e= "2'-methoxyethyl (2'-MOE) nucleotides"
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Pred. No. 8.7e+02;
0; Mismatches ]
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RESULT 745
ABQ93238
ID ABQ932
XX ABQ932
XX 29-AUG
DT 29-AUG
DT 21-CCT
XX Of the Micros
KW Micros
KW Micros
KW Giseat
KW resiat
KW geneti
XX Pellon
CX Aegilo
OS Tritic
XX Aegilo
OS Tritic
XX EP1217
XX 22-DEC
XX Map of
PT Map of
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MicroBatellite marker; wheat; D genome; mapping; genotyping; polymorphism; phenotypic trait; QTL; quantitative trait locu disease-associated gene; development factor; quality factor; resistance factor; wheat product; identification; detection; genetically modified wheat; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to antisense compounds which inhibit the express of human caspase 8. The antisense compound is useful for diagnosting at treating diseases associated with the expression of caspase 8 and for prophylaxis e.g. to prevent or delay infection, inflammation or tumous formation, and as a research reagent. The present sequence is an
                                                                                                                                                                                                                                                                                     Map of wheat D genome comprising the genome location of a microsatellite marker, useful for e.g. identifying genes responsible for a desired phenotypic trait, especially quantitative trait loci in wheat, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New antisense compounds capable of modulating expression of caspase the diagnoses, prophylaxis and treatment of diseases associated with expression of caspase 8, e.g. inflammation and tumor formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
                                                                                                                                                                                                       Claim 4; Page 8; 105pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bernard
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22-DEC-2000; 2000EP-00403659
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21-OCT-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sourdille P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (revised)
(first en
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Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    15.4; DB 1;
No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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The invention relates to a map of the bread wheat D genome comprising the genome location of a microsatellite marker selected from a group of 185 such markers (ABQ92733-ABQ92917). The invention also encompasses the use of left (ABQ92703-ABQ93102) and right (ABQ93103-ABQ93287) primers to amplify and detect the microsatellite markers, and to identify genes responsible for a phenotypic trait of interest in wheat. Wheat is an

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RESULT 746
AAD34340
ID AAD343
XX AAD343
XX I6-JUL
XX Human
DE Human
EW Human;
KW Human;
KW bone d
KW bone d
KW therap;
KW bone d
KW therap;
KW osteop
XX Homo &
XX Homo &
XX I6-AUC
XX Homo &
XX I6-AUC
PN WO2002
XX PN WO2002
XX PN WO2002
XX I6-AUC
PR 18-AUC
PR 18-AUC
PR 18-AUC
PR 18-AUC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC allohexaploid species consisting of 3 diploid genomes designated A, B and CD, resulting from two successive intercrossing involving at least three cdifferent species. The D genome is thought to have been introduced in the CC most recent intercrossing, between the amphiploid AABB and Triticum CC tauschii (DD), probably involving only a limited number of genotypes of CC both species. Due to its polyploid genome, the large size of its genome, CC date been difficult. Microsatellites are tandemly repeated sequences and its low level of polymorphism, the genetic mapping of wheat has to CC date been difficult. Microsatellites are tandemly repeated sequences cc between one and six nucleotides long, and are very polymorphic in length, CC mainly due to polymerase slippage during replication. This high degree of CC polymorphism makes them especially suitable for the genetic mapping of CC species which show little intraspecies polymorphism, such as wheat. In CC distance. The 185 microsatellite markers of the invention are useful for identifying donor species Triticum tauschii and CC map to the wheat D genome, which is less polymorphic than the A or B CC groomes. These microsatellite markers bus help to overcome some of the CC map and the microsatellite markers and associated primers of the CC invention are useful for identifying genes responsible for a phenotypic CC trait of interest, most notably QTLs (quantitative trait loci). In CC disease and for identifying development factors, quality factors and CC factors conferring resistance to pathogens and an allelse implicated in CC mapping and genotyping diploid and polyploid species of Triticum, CC mapping and genotyping diploid and polyploid species of Triticum, CC mapping and genotyping diploid and polyploid species of Triticum durum, Triticum account, Triticum durum, Triticum account and polyploid species and to assess whether or not a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                   bone strength; mineralisation; ophthalmological; antidiabetic; bone density regulating transmembrane receptor; prosthetic device; surgical implant; diabetic retinopathy; hypertensive retinopathy; therapy; osteoporosis; prematurity; ocular vessel; eye disorder; osteopathic; PCR; primer; ss.
      18-AUG-2000;
22-SEP-2000;
                                                                                                  17-AUG-2001; 2001WO-US041788
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; bone strength and mineralisation regulatory protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human BSMR gene polymorphism detecting PCR primer, LRGEN9R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16-JUL-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAD34340 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3423 GAGGCAGGAACTGAGGG 3439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 6 A; 2 C; 10 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
      2000US-0226119P
2000US-0234337P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BSMR;
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RESULT 747
ABX64386/c
   PRRACE NO CONTRACTOR NO CONTRA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to bone strength and mineralisation regulatory composed in (BSWR) and its corresponding nucleic acid sequence. BSWR DNA is consequence in the diagnosis or therapy of osteoporosis and for regulating continuous plants or therapy of osteoporosis and for regulating continuous plants of the diagnosis or therapy of osteoporosis and for regulating continuous plants of the diagnosis or therapy of osteoporosis and for regulating a promoter that is operably continuous plants of the diagnosis of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local (
11-JUN-1993;
10-JUN-1994;
20-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR; primer; ss; retroviral packaging plasmid; retrovirus; gene gene therapy; replication-incompetent; replication-competent; helper virus; LTR; long terminal repeat; virus; psi; xenotropic; amphotropic; ecotropic; polytropic; envelope protein; haematopoi stem cell; lymphocyte.
                                                                                                                                                                                                                                                                                                                                                    Gibbon Ape Leukemia Virus.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABX64386;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20 BP; 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Fig 5; 124pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Polynucleotide encoding bone strength and mineralization regulatory protein useful for diagnosis or therapy of osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Warman ML, Gong Y,
                                                                                                                                           04-SEP-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Retroviral packaging plasmid construction PCR primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-AUG-2003
28-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABX64386 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              13-JUL-2001; 2001US-0304851P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (AVET ) AVENTIS PHARMA SA. (HARD ) HARVARD COLLEGE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (UYCA-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3422 TGAGGCAGGAACTGAGG 3438
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                                                                                                                                              2001US-00944411.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (revised)
(first entry)
93US-00076299.
94US-00258152.
97US-00914893.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       з c;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Olsen BR, Rawadi G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
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Pred. No. 8.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       U; 0 Other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein; haematopoietic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     gene delivery;
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ACC45221/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PINE/) FINER M H.
(DULL/) DULL T J.
(ZSEB/) ZSEBO K M.
(COOK/) COOKB K.
(FARS/) FARSON D A.
                                                                                                              Human; cytostatic; nootropic; neuroprotective; antiinflammatory; antisense therapy; NAC; DEFCAP; hyperproliferative disease; apoptosis; death effector filament-forming CED4-like apoptosis protein; neurological disease; infection; inflammation; tumour formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Novel retroviral packaging system useful for producing high-titers of recombinant retrovirus in human cells, comprises at least one retrovi helper DNA sequence derived from replication-incompetent retroviral
                                                                                                                                                                                                                                                16-JUN-2003
                                                                                                                                                                                                                                                                                                                         ACC45221 standard; DNA; 20
                                   Synthetic.
                                                         Homo sapiens
                                                                                                                                                                                                         Human NAC chimeric phosphorothioate oligonucleotide SEQ ID NO:81.
                                                                                                                                                                                                                                                                                      ACC45221;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 2 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 14; Page 27; 53pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Finer MH,
                                                                                            phosphorothicate;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        4821 CACCAGCCCTTGACCCT 4837
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16;
                                                                                                                                                                                                                                                                                                                                                                                                                    19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                    CACCAGCCCATGACCCT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                              (first entry)
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.3%;
94.1%;
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                                                                                                                                                                                                                                                                                                                         ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <u>;</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.4; DB 1; Length 20; Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cooke K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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ABT43248
ID ABT4
XX
AC ABT4

ABT43248 standard; DNA; 20

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ABT43248;

RESULT 749

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cc at least an 8-nucleobase portion of an active site on a nucleobase portion of an active site on a nucleobase portion of an active site on a nucleobase portion of an active site on a nucleoic acid composition (I) cand a pharmaceutical carrier or diluent; (2) inhibiting the expression of CC NAC in cells or tissues comprising contacting the cells or tissues with CC (I); and (3) treating an animal having a disease or condition associated with NAC comprising administering (I) to the animal so that expression of CC NAC is inhibited. (I) has cytostatic, nootropic, neuroprotective and CC antisnflammatory activities, and can be used in antisense therapy. The cantisense compounds (I) are useful for modulating the expression of NAC, C and for treating a disease or condition associated with expression of CC and for treating a disease, neurological disease, or a disease or disorder arising from aberrant apoptosis. The compounds are also useful as research reagents and kits, or for disgnostics, therapeutics cuseful as research reagents and kits, or for disgnostic, therapeutics compounds are also known as a death effector filament-forming CC Lumour formation. NAC is also known as a death effector formation or this protein (DEPCAP). NAC is located on human chromosome CC phosphorothicate antisense oligonucleotide, which is given in the present invention.
                                                                                                                                                          Query Match
Best Local &
                                                                                                                           Matches
                                                                                                                                                                                                                                                          Sequence 20 BP; 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a compound (1) 8-50 nucleobases in length targeted to a nucleic acid molecule encoding NAC, where the compound specifically hybridises with the nucleic acid molecule encoding NAC and inhibits the expression of NAC. The compound specifically hybridises with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 15; Page 76; 147pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compounds, useful for modulating the expression of Ni for treating a disease or condition associated with the expression NAC, e.g. hyperproliferative disease or neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        19-SEP-2001; 2001US-00956712.
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                                                                                                                                                                                                                                                                                                                        exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-SEP-2002; 2002WO-US029664.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-) ISIS PHARM INC
                                                                                                                           Local Similarity
nes 16; Conserv
                                                          4303 AAGGAACTGGAGCTCTG 4319
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AAGGAACTGGAGCTTTG 1
                                                                                                                        Conservative
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16. .20
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/note= "2'-O-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/mod_base= OTHER
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                                                                                                                                                                                                                                                          A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                      0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "2'-O-methoxyethyl (2'-MOE) gapmer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ი
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                                                                                                                           0;
                                                                                                                                                          Score 15.4; DB 1;
Pred. No. 8.7e+02;
                                                                                                                              Mismatches
                                                                                                                                                                                         Length 20;
                                                                                                                                  Indels
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                                                                                                                              Gaps
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RESULT 750
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic acids isolated from neuroblastoma showing enhanced expression in human neuroblastoma with good prognosis, useful in clarifying good/poor prognosis of neuroblastoma and providing genetic data.
                                                                                30-MAY-2001; 2001JP-00162775.
24-AUG-2001; 2001JP-00255226.
                                                                                                                                                            30-MAY-2002;
                                                                                                                                                                                                               05-DEC-2002.
                                                                                                                                                                                                                                                                                                                      Unidentified.
                                                                                                                                                                                                                                                                                                                                                                 high malignancy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Neuroblastoma-related oligonucleotide #196.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ABT32419 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention comprises DNA sequences that show enhanced expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nakagawara
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24-AUG-2001; 2001JP-00255260.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Neuroblastoma-related DNA
                                                                                                                                                                                                                                                                WO200297093-A1
                                                                                                                                                                                                                                                                                                                                                                                         Neuroblastoma; prognosis; spontaneous regression; primer; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 5; Page 24; 444pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2003-167523/16.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human neuroblastoma with good prognosis. The DNA sequences of the invention are useful in clarifying good/poor prognosis of neuroblastoma. The present DNA sequence was used in the exemplification of the inventic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HISM ) HISAMITSU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
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  CHIBA PREFECTURE.
HISAMITSU PHARM CO LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; G protein-coupled receptor; hyperproliferative disorder; GPR37L1; endothelin type b receptor-like protein-2; cerebral vascular disease; antisense; endothelin-binding receptor-like protein-2; atherosclerosis; cardiovascular disease; ETBR-LP-2; G-protein coupled receptor 37 like 1; acute proliferative nephropathy; ETBR-like protein 2; cancer; stroke; angiogenesis; hypertension; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention comprises nucleic acids that show increased expression in human neuroblastomas with poor prognosis over those with a good prognosis. The nucleic acids of the invention are useful as a tool for distinguishing neuroblastomas with a favourable prognosis (spontaneous regression) from neuroblastomas with a poor prognosis (high malignancy). The DNA sequences ABT32224 - ABT32571 represent oligonucleotides used in an example of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 8 A; 6 C; 4 G; 2 T; 0 U; 0 Other;
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                                                   06-DEC-2001; 2001US-00003126.
                                                                                                       04-DEC-2002; 2002WO-US038520.
                                                                                                                                                             19-JUN-2003.
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(ISIS-) ISIS PHARM INC
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are 5-methylcytidines"
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Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                     (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                          (2'-MOE)
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RESULT 752
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Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to antisense compounds targetted to the nucleic acid encoding the G protein-coupled receptor ETBR-LP-2 (endothelin type b receptor-like protein-2) to inhibit its expression. ETBR-LP-2 is also known as endothelin-binding receptor-like protein-2, ETBR-LP-2 is also and G-protein coupled receptor 37 like 1 (GPR37L1). Antisense compounds of the invention are useful for treating hyperproliferative disorders (especially cancer) and cardiovascular diseases especially angiogenesis, atherosols, hypertension, cerebral vascular disease, stroke and acute proliferative nephropathy. The present sequence is an antisense oligonucleotide targetted to human ETBR-LP-2 DNA
The invention relates to screening for an oestrogen analogue, comprising administering a test compound to rodents, isolating hippocampus from rodents, monitoring for the expression level of a gene comprising mouse calmodulin I, chimaerin, neuromedin, T-Rec-alpha rel., 3B-HSD related protein, ATP-binding cass., chaperonin, HCNP, histone-like protein, unknown, ZW10, vitronectin or unknown encoding genes (SEQ ID NO 1-13) in
                                                                                                                                                                Screening for estrogen analog, by administering test compound to rodents, isolating hippocampus, monitoring for the expression of a particular gene in hippocampus, and selecting compound that alters gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mouse; oestrogen; hippocampus; gene expression; calmodulin I; chimaerin; neuromedin; T-Rec-alpha rel.; 3B-HSD related protein; ATP-binding cass.; chaperonin; HCNP; histone-like protein; ZW10; vitronectin; gene; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New oligonucleotides which bind the nucleic acid encoding the G protein coupled receptor ETBR-LP-2 (endothelin type b receptor-like protein-2 receptor), useful for treating e.g. cancer and cardiovascular diseases.
                                                                                                                                   Disclosure; Fig
                                                                                                                                                                                                                                                                                                                           02-NOV-2001, 2001JP-00338515.
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nilarity 94.1%;
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                                                                                                                                 2; 16pp; Japanese.
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Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Caspase-8; cytostatic; immunosuppressant; anti-HIV; ss; antisense gene therapy; apoptosis; hyperproliferative disorder; haematopotetic disorder; autoimmune disorder; viral infection; A neurological disorder; Alzheimer's disease; Parkinson's disease; amyotrophic lateral sclerosis; retinitis pigmentosa; blood cell
                                                                                                                                           WPI; 2003-810793/76.
                                                                                                                                                                                                                                      (ZHAN/)
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11-JAN-2001; 2001WO-US000955
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| 16; Conserv
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COWSERT L M.
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No. 8.7e+02;
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New compounds, particularly antisense oligonuclectides targeted to a nucleic acid encoding caspase 8, useful for treating a disease/condition associated with caspase 8, such as hyperproliferative or autoimmune

disorders.

Claim

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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Matches 16
                                                                                  Detecting a variant in a primer extension assay, useful for analyzing molecular events for identifying mutations indicative of cancer, by contacting a target nucleic acid primer complementary to a region of trarget nucleic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                 variant detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Variant detecting primer extension wild-type product oligo, SEQ ID
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                                                                                                                                                                                                                                                                 15-FEB-2002; 2002US-0357585P
                                                                                                                                                                                                                                                                                                 18-FEB-2003; 2003WO-US004827.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 caspase 8, i.e. an antisense oligonucleotide. Also included 8-30 nucleobases in length that specifically hybridises
                                                                                                                                                                                                                                                                                                                                                                                                                                ion; primer extension assay; mutation; cancer;
sporadic mutation; genotyping; pooled sample;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NO 168; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.3%;
                                                                                                                                                                                                 Whitney
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Pred.
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The invention relates to a novel method for detecting a variant primer extension assay, useful for analysing molecular events for

Claim 15; SEQ ID NO 913; 872pp; English

corresponding RNAs,

The invention relates to

a

novel pharmaceutical

composition, which has

Example 7;

SEQ ID NO 110; 54pp; English

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RESULT 755
ABZ85671/C
IID ABZ856
XX ABZ856
XX ABZ856
XX ABZ856
XX Human
XX Human
XX Human
XX Human
XX Homo 8
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Detecting a variant in a primer extension assay comprises contacting a target nucleic acid primer complementary to a region of the target contacting a complementary to a region of the target nucleic acid, and extending the primer in the presence of a first conclectide that is complementary to a first variant nucleotide suspected to be at a position downstream of the region and a second nucleotide that is complementary to a second variant nucleotide at the position, thus to reduce misincorporation of the first nucleotide on a template comprising the second variant nucleotide. The methods are useful for analysing molecular events for identifying individuals with mutations indicative of cancer. They are particularly useful in detecting a rare mutation in a heterogeneous biological sample (e.g. sporadic mutation in a heterogeneous patient sample), detecting rare genotypes in genotyping reactions; or detecting mutant or viral sequence variations in pooled patient samples). This polynucleotide sequence represents an oligo used as part of the primer extension assay
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antianflammatory; steroid; ubiquinone; entinflammatory; antialized; hypotensive; immunosuppressive; cytostatic; gene therapy antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                        Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human oligonucleotide sequence
                                                                                                                                                                                                                                                                                   WPI; 2003-229219/22.
                                                                                                                                                                                                                                                                                                                                             Miller
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                                                                                                                                                                                                                                                                                                                                                                                                                                           (EPIG-)
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                                                                                                                                                                                                                                                                                                                                       Li Y, Sa
Tang L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 3 A; 4 C; 9 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                              Sandrasagra A,
                                                                                                                                                                                                                                                                                                                                                                                                                                           PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         respiratory disease; ds.
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                                                                                                                                                                                                                                                                                                                                                    Shahabuddin
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Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                                 Pabalan J,
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RRSULT 756
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                     Nyce JW,
Miller S,
                                                                                                                                                                   Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; antisense; lung dysfunction; nasal airway dysfunction;
antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    31-OCT-2002.
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antisense gene therapy; respiratory; lung; adenosine sensitivity;
adenosine receptor; bronchodilation; bronchoconstriction; lung allergy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human oligonucleotide sequence.
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                                                                                                                                                                                                                                                                                                        WPI; 2003-229219/22.
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                                                                                                                                                                                                                                                                                                                                                                  Li Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                     Sandrasagra A,
C, Shahabuddin
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Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disease; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                    Katz E,
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The invention relates to a novel pharmaceutical composition, which has

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Claim 15; SEQ

ID NO 1314; 872pp; English

invention relates to a novel pharmaceutical composition, which has

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Disclosure; SEQ ID NO 3824; 872pp; English

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RESULT 757
ABZ86072
ID ABZ8607
XX ABZ860
XX ABZ860
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DT 17-OCT
XX Human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                  Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23-APR-2002; 2002WO-US013135
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                                                                                                                                                                              corresponding RNAs,
                                                                                                                                                                                                                                                                                                                                      WPI; 2003-229219/22
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, Tang L,
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                                                                                                                                                                                                                                                                                                                                                                                                       Sandrasagra A,
L, Shahabuddin
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Pred. No. 8.7e+02;
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RESULT 758
ABD21901/c
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XX Human
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KW respir
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; antiinflammatory; antisethmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstriction; respiratory distress syndrome; allergic rhinitis; pulmonary hypertension; emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
                                                                                  Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
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5' and 3' intron-exon junctions,
                                                                                                                                                                                                                                                                                                                                                                                           24-APR-2001; 2001US-0286036P
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Tang
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L, Shahabuddin
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Pred. No. 8.7e
0; Mismatches
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ಸಾಗ. 8.7e+02;
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comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and creducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, composition and depletion or hyposecretion, when administered to a mammal. The oligonucleotides are derived from a gene encoding or regulating composition or cancer and can be anti-sense to the corresponding mRNA. The invention also describes a ktt, that comprises: (a) a delivery composition of a target polypeptide associated with lung airway or lung daysfunction or cancer and can be anti-sense to the corresponding mRNA. The invention has antiallergic, antinflammatory, antiasthmatic, analysesic, hypotensive, immunosuppressive and cytostatic activity, is a beta-adrenergic agonist. The composition of the kit. The composition of the invention of the composition of the comprises oligo and is administered to reduce the production corresponding are respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The composition comprises oligo and is administered to reduce the production, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, 
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Best Local Similarity
Matches 16; Conserv
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ABD24812 standard; DNA; 20
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                                                                                                                                                                                                                                                                Score 15.4; DB 1;
Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                            Length
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ABD24812
ID ABD248
XX ABD2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AI092623-derived oligonucleotide SEQ ID 3824.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pulmonary transplantation rejection; ss; primer.
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bronchodilating

agent. NO 913;

> 24-APR-2001; 2001US-0286036P. 23-APR-2002; 2002WO-US013143

SEQ ID

763pp; English

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RESULT 760
ABD22302
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         This invention describes a novel composition (a) a first active agent, comprising oligonucleotides, effective for alleviating comprising oligonucleotides, effective for alleviating allergies and crediting adenosine sensitivity, levels of adenosine (A) or (A) receptors, comprising adenosine sensitivity, levels of adenosine (A) or (A) receptors, comprises on of a target polypeptide associated with lung airway or lung compression of a target polypeptide associated with lung airway or lung compression of a target polypeptide associated with lung airway or lung compression of a target polypeptide associated with lung airway or lung compression of a delivery (C) the invention also describes a kit, that comprises: (a) a delivery (C) device, in separate containers, (b) the oligonucleotides, (c) analgesic, hypotensive, immunosuppressive and cytostatic activity, is a compression for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiallammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a compressive, immunosuppressive and cytostatic activity, is a composition of the composition of the target manual compressive and analgesic activity, is a composition of the target manual composition are advantable of composition, and/or malignant disease. The administered composition, and/or bronchoconstriction and/or lung unmanual compresses and compresses the degradation of the target manual composition, and/or bronchoconstriction and/or lung unmanual composition, and/or bronchoconstriction and/or lung unmanual compression, and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vasoconstriction, compresses and composition of the anti-sense oligon corresponding to the anti-sense oligon corresponding to the anti-sense oligon corresponding to the composition of the system of the composition of the breakdown of the composition of the composition of the system of the composition of the composition of the system of 
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Best Local (
Human; antisense; bronchoconstriction; allergy; hyposecretion; pain; respiratory tract inflammation; adenosine sensitivity; lung; cancer; surfactant depletion; antiallergic; artiinflammatory; antiasthmatic; analgesic; hypotensive; immunosuppressive; cytostatic; cystic fibrosis; beta-adrenergic agonist; respiratory disease; pulmonary vasoconstrictio respiratory distress syndrome; allergic rhinitis; pulmonary hypertensio emphysema; chronic obstructive pulmonary disease; cancer; bronchitis; pulmonary transplantation rejection; ss; primer.
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Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic golds associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                              Human stanniocalcin-derived oligo SEQ ID 1314.
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Tang
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                           (first
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L, Shahabuddin S;
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94.1%;
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0; Mismatches
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Pred. No. 8.7e+02;
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                                                                           pulmonary vasoconstriction;
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RESULT 761
ADH08330/c
ID ADH083
XX
AC ADH083
XX

ADH08330 standard;

DNA;

20

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ADH08330

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2646 GCTGCTGCTGCAGCCAC 2662

Similarity

Conservative

0;

Mismatches

Gaps

0

8.7e+02

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N

18

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CC compositing oligonucleotides, effective for alleviating composition, respiratory tract inflammation, allergies and cc reducing adenosine sensitivity, levels of adenosine (A) or (A) receptors, cc surfactant depletion or hyposecretion, when administered to a mammal. The coligonucleotides are derived from a gene encoding or regulating cc expression of a target polypeptide associated with lung airway or lung cyperate in the invention also describes a kit, that comprises: (a) a delivery cc instructions for adding a carrier and for use of the kit. The composition of the invention has antiallergic, antiinflammatory, antiasthmatic, analgesic, hypotensive, immunosuppressive and cytostatic activity, is a cc treating a respiratory, lung or malignant disease. The administered composition comprises oligo and is administered to reduce the production or availability, or to increase the degradation of the target mRNA or to reduce the amount of target polypeptide present in the lungs. The cc pulmonary obstruction, and/or bronchoconstriction and/or lung inflammation, allergies and/or surfactant hypoproduction are associated with a disease or condition such as pulmonary vascoonstriction are associated conflammation, emphysema, chronic obstructive pulmonary disease, pulmonary transplantation rejection, pulmonary infections, bronchitis or cancer. The reduced adenosine content of the anti-sense oligos corresponding to thymidines present in the target RNA serves to prevent the breakdown of the oligonucleotides into products that free adenosine into the system ce.g., lung, brain, heart, kidney, etc, tissue environment and thereby, to prevent any unwanted effects due to it
Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nyce JW,
Miller S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pharmaceutical composition for treating asthma, has antisense oligonucleotide containing less percentage of adenosine, targeted to nucleic acids associated with lung airway or lung dysfunction, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 15; SEQ ID NO 1314; 763pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-APR-2001; 2001US-0286036P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               23-APR-2002; 2002WO-US013143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200285309-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Li Y,
Tang
                                                                                                                                                 BP; 1 A; 8 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      describes a novel composition (a) a first active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sandrasagra A,
L, Shahabuddin
                              0.3%;
                                      Score 15.4;
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ø
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                                                                        DB 1; Length 20;
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ARBSULT 762
ADH12202
ID ADH122
AC ADH122
AC ADH122
XX
PT 11-MAR
XX
UMAR
DE Human;
KW Human;
KW Chroma
KW Human;
KW Chroma
KW Gevelo
KW develo
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to a biochip (1) that comprises a support (2) the functional side of which has a working surface (3) with a network of elementary sites (Xn), with many ligands, different for each Xn, attached to them. The new feature is that Xn are distributed between: a central zone (4), designed for detection of at least one target species, with each site containing ligands for a particular target; and a peripheral zone (5), surrounding (4) and containing control sites, which optionally carry control ligands. The biochip is used for analysis in molecular biology. The specified arrangement of sites allows not only determination of many targets but also monitoring of the determination and of the operating conditions. The signals emitted from the chip are not affected by the geometrical environment of the chip, for any of the elementary sites. The current sequence represents a fragment of a mutant gene that is related to career
                                                   Human; chromodomain helicase DNA-binding 5; CHD5; chromosome 1p36.3; chromatin structure; chromatin unwinding; DNA repair; DNA recombination; transcriptional regulation; gene expression; cell cycle control; development regulation; oncogenesis; brain; neural development; neural tissue neoplasia; diagnosis; cancer; neural cancer; neuroblastoma; breast cancer; colon cancer; liver tumour; germ cell tumour;
                                                                                                                                                                                   Human CHD5 PCR/sequencing primer,
                                                                                                                                                                                                                       11-MAR-2004 (first entry)
                                                                                                                                                                                                                                                             ADH12202
                                                                                                                                                                                                                                                                                               ADH12202 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biochip, useful in molecular biology, comprises a central array of sites carrying analytical ligands and a peripheral region containing control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cuzin M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          24-MAY-2002; 2002FR-00007039
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            23-MAY-2003; 2003WO-FR001574
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Biochip; target; ligand; analysis; molecular biology;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    11-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            related to cancer
                                                                                                                                                                                                                                                                                                                                                                                                                           4737 GGAGACCCATCCTCACC 4753
                                      screening;
                                                                                                                                                                                                                                                                                                                                                                                         20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Page 17;
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                                                                                                                                                                                                                                                                                                                                                                                         GGAGGCCCATCCTCACC 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                      cytostatic; gene therapy; sequencing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; 4 C; 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cleuziat P,
                                                                                                                                                                                                                                                                                               ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15.4; DB 1;
Pred. No. 8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                   SEQ ID NO:24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                      primer; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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The invention relates to human chromodomain, helicase, DNA-binding 5 (CHD5; ADH12180) and cDNA encoding it (ADH12179). CHD5 is a novel member CC (chromatin structure and gene expression and which have a profound effect on CC chromatin structure and gene expression and which are thus likely to play CC an important role in cell cycle control, regulation of development, and CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central region which CC (plant homeodomain) class, two chromodomains, a central regulators of the PHD (plant homeodomain) class, and chastify deleted in a variety contains or crelates to vectors and host cells comprising the CHD5 cDNA sequence; call very correct or cancer; treating cancer by administration of CHD5 protein, CHD5 encoding CC (plant homeodomains) creates such as neural cancers (e.g., neuroblastoma), breast cancer, colon cancers such as neural cancers (e.g., neuroblastoma), breast cancer, CHD5 mucleic acids and anti-CHD5 antibodies are useful as comprised or chromatin unwinding, DNA repair and recombination, and transcriptional crematation. Sequences ADH12186-ADH12247 represent specifically claimed components.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003106650-A2
                                               regulation. Sec
human CHD5 PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diagnostic and prognostic indicator of tumor, comprises amino thaving two PHD class zinc finger domains and two chromodomains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel chromodomain helicase DNA-binding (CHD) proteins, useful as diagnostic and prognostic indicator of tumor, comprises amino terminus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2004-082186/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brodeur GM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-JUN-2002; 2002US-0388848P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-JUN-2003; 2003WO-US019027.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CHILDRENS HOSPITAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 24; 124pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             White PS;
                                               primers.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PHILADELPHIA
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Sequence 20 BP; 7 A; 5 C; 5 G; 3 T; 0 U; 0 Other;

Matches Query Match Best Local 16; Similarity Conservative 0.3%; 0; Score 15.4; DB 1; Pred. No. 8.7e+02; Mismatches Length 20; Indels 0, Gaps

0

3313 CAGAACAACCTGGATGA 3329 CAGAACAACCTGGAGGA 20

밁 8

ADJ45268 standard; DNA; 20 ВЪ

ADJ45268;

06-MAY-2004 (first entry)

Hepatoma-derived growth factor antisense oligo seqid 38.

RESULT 763
ADJ45268
ID ADJ452
XX ADJ452
XX ADJ452
XX O6-MA)
XX XX
DT 06-MA)
XX XX
XX Cytost
XX Cytost
XM hepatc
XM hepatc
XM humanı cytostatic; endocrine; hepatoma-derived growth factor inhibitor; hepatoma-derived growth factor; metabolic disorder; hyperprolife oligonucleotide disorder; hyperproliferative;

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                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                 Query Match
Best Local
                                                                                                                                                                                                                                                                                             useful for treating a disease or condition associated with hepatoma-derived growth factor, such as a metabolic disorder, or a hyperproliferative disorder, e.g. cancer, which is selected from hepatoma, leiomyoma, esophageal cancer or ovarian cancer. They are also useful in research and diagnostics for modulating the expression of hepatoma-derived growth factor. This sequence represents a human hepatom-derived growth factor antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                      The invention describes a compound 8-80 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding hepatoma-derived growth factor, and inhibits the expression of hepatoma-derived growth factor. The compound, composition and methods are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New compound, particularly an antisense oligonucleotide targeted to nucleic acid encoding a hepatoma-derived growth factor, useful for treating a hyperproliferative disorder e.g. cancer, or a metabolic
 cytostatic; endocrine; hepatoma-derived growth factor inhibitor;
                                                    06-MAY-2004
                                                                             ADJ45339;
                                                                                                       ADJ45339
                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 15; SEQ ID NO 38;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-142660/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-JUL-2002; 2002US-00210429.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-JUL-2002, 2002US-00210429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-FEB-2004.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                         Hepatoma-derived growth factor antisense oligo seqid 109.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ISIS-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  modified_base
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                                                                                                                                                                                          1103 TAGCACCCAGAGAGCAG
                                                                                                                                                                    4
                                                                                                                                                                                                                      16;
                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ISIS PHARM INC.
                                                                                                     standard;
                                                                                                                                                                    TAGCACCCAGACAGCAG
                                                                                                                                                                                                                                                                        BP;
                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dobie KW;
                                                    (first entry)
                                                                                                                                                                                                                                                                      7 A; 7 C; 5 G; 1 T; 0 U;
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15. .20
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/mod_base= OTHER
/note= "OTHER= 2'-O-Methoxyethyl (2'-MOE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /mod_base= OTHER
/note= "OTHER= Phosphorothioate backbone. All cytidines
/note= "OTHER= Phosphorothioate backbone. All cytidines"
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                                                                                                     DNA;
                                                                                                                                                                                                                               0.3%;
94.1%;
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e= "OTHER= 2'-O-Methoxyethyl (2'-MOE)
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                                                                                                       80
                                                                                                                                                                                                                                                                                                                                                                                                                                                      61pp; English.
                                                                                                                                                                                           1119
                                                                                                                                                                    20
                                                                                                                                                                                                                     <u>;</u>
                                                                                                                                                                                                                                 Score 15.4;
Pred. No. 8.
                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                              8.7e+02
                                                                                                                                                                                                                                                                      0 Other;
                                                                                                                                                                                                                                             DB 1; Length 20;
                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               nucleotides'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleotides"
                                                                                                                                                                                                                     0
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1 hepatoma
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ADL59589
ID ADL5
XX
AC ADL5
XX
DT 03-0
                                                                                                                                                                            RESULT 765
                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention describes a compound 8-80 nucleobases in length targeted to, and which specifically hybridises with a nucleic acid molecule encoding hepatoma derived growth factor. The compound, composition and methods ar useful for treating a disease or condition associated with hepatoma-derived growth factor, such as a metabolic disorder, or a hyperproliferative disorder, e.g. cancer, which is selected from hepatoma, leiomyoma, esophageal cancer or ovarian cancer. They are also useful in research and diagnostics for modulating the expression of hepatoma-derived growth factor. This sequence represents a human hepatom derived growth factor antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New compound, particularly an antisense oligonucleotide targeted nucleic acid encoding a hepatoma-derived growth factor, useful for treating a hyperproliferative disorder e.g. cancer, or a metaboli
  Human ESM-1 antisense oligonucleotide segid 1838.
                                             03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 1 A; 5 C; 7 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      disorder
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bennett CF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-JUL-2002; 2002US-00210429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        31-JUL-2002; 2002US-00210429.
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                                                                                                                                   ADL59589 standard;
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/mod_base= OTHER
/mote= "OTHER= 2'-0-Methoxyethy1
/note= "OTHER= 2'-0-Methoxyethy1
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are 5-methylcytidines"
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note= "OTHER= 2'-O-Methoxyethyl (2'-MOE)
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Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                                                                                                                     Length 20;
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ADL59612
                               ADL59612 standard; DNA; 20 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
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                                                                                                                                     2892 CCCAGTCACAGATGCCT 2908
                                                                                                                                                                                                                                                                                                                                                                                                                                          3; SEQ ID NO 1838; 555pp; English.
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/note= "OTHER= 2
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94.1%;
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                                                                                                                                                                              Score 15.4; DB 1;
Pred. No. 8.7e+02;
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                                                                                                                                                                                                                                                                           The invention describes a new antisense compound, having a sequence comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition; inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury, or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonucleotide that can be used to modulate expression of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
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                                                                                                                                                                                                  Sequence 20
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                                                                                                                                                                                                                                                        specific molecule-1 (ESM-1).
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                                                                                                     Conservative
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16. .20
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/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
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                                                                                                          Sequence 20
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ilarity 94.1%;
Conservative
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/note= "OTHER= 2
16. .20
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/note= "OTHER= phosphorothioate backbone. All cytidine
residues are 5-methylcytidines"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                             9 C; 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       base= OTHER
== "OTHER= 2'-methoxyethyl (2'-MOB) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    555pp, English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP
0,
                                                                                                             G; 5 T; 0 U; 0 Other;
  Pred. No. 8.7
); Mismatches
                            Score 15.4;
Pred. No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2'-methoxyethyl (2'-MOE)
                       8.7e+02
                                                      DB 1; Length 20;
     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nucleotides"
  0
  Gaps
     0
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S

2892 CCCAGTCACAGATGCCT 2908

Query Match Best Local Similarity

0.3%; 94.1%;

Score 15.4; DB 1; Pred. No. 8.7e+02;

Length

20

Sequence 20

BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;

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RESULT 768
ADL59695
ID ADL596
XX ADL596
XX ADL596
XX Gytost
KW gene t
KW gene t
KW gene t
KW gene t
KW Gytost
KW gene t
KW Gytost
KW gene t
KW Gytost
FT Hodifi
FT modifi
FT mod
                                                  comprising 8-30 bp targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), that specifically hybridises with the nucleic acid ESM-1 and inhibits its expression. Also described are: a composition, inhibiting the expression of ESM-1 in cells or tissues; and treating an animal having a disease or condition associated with ESM-1. The compound is useful for preparing a composition for treating diabetes, cancer, ischaemia or reperfusion injury or angiogenic, immunological, cardiovascular or neurological disorder. This sequence represents an antisense oligonucleotide that can be used to modulate expression of endothelial specific molecule-1 (ESM-1).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New antisense compound, having a sequence targeted to a nucleic acid encoding endothelial specific molecule-1 (ESM-1), useful for preparing composition for treating e.g., diabetes, cancer or cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cytostatic; antidiabetic; immunomodulator; cardiant; neuroprotective; gene therapy; endothelial specific molecule-1; ESM-1; ESM-1 ESM-1 ralated disorder; diabetes; cancer; lachaemia; reperfusion injury; angiogenic disorder; immunological disorder; cardiovascular disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human
                                                                                                                                                                                                                                                                                                                                                                   The invention describes a new antisense compound, having a sequence
                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 3; SEQ ID NO 1944; 555pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19-AUG-2002; 2002US-0404495P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-AUG-2003; 2003WO-US025833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           neurological disorder; antisense technology; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PHAA ) PHARMACIA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 CCCTGTCACAGATGCCT
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/note= "OTHER= phosphorothioate
residues are 5-methylcytidines"
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/mod_ba
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e= "OTHER= 2'-methoxyethyl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ი
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RESULT 769
AD0425
AD0425
XX AD0426
XX AD0425
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    11-DEC-
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05-DEC-2001;
07-DEC-2001;
07-DEC-2001;
10-DEC-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; NOVX; PCR; ss; cancer; atherosclerosis; diabetes; Alzheimer's disease; Parkinson's disease; graft-versus-host disease scleroderma; hypertension; happendphila; idiopathic thrombocytopenic purpura; immunodeficiency; AIDS; dyslipidemia; obesity; Crohn's disease; bronchial asthma; anorexia; cancer-associated cachexia; multiple sclerosis; fertility; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               02-DEC-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2004
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US2004058338-A1
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                                                                   1-2002;
1-2002;
1-2002;
1-2002;
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1-2002;
1-2002;
1-2002;
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-2001;
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-2001
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2001US-0338285P
2001US-0338318P
2001US-0339314P
2001US-0339517P
2001US-0339517P
2001US-0349611P
2001US-0349611P
2001US-0349640
2001US-03496608P
2001US-03411440P
2001US-0341147P
2001US-0341147P
2001US-0341178P
2001US-0341540P
2001US-0341540P
2001US-0341540P
2001US-034592P
2002US-0359671P
2002US-0359671P
2002US-0359671P
2002US-0359671P
2002US-0361264P
2002US-0361264P
2002US-0361264P
2002US-036230P
2002US-036230P
2002US-036239P
2002US-036239P
2002US-036239P
2002US-036239P
2002US-0364238P
2002US-0365025P
2002US-0365025P
2002US-0365025P
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         15-MAY-2002;
11-MAY-2002;
17-MAY-2002;
28-MAY-2002;
28-MAY-2002;
29-MAY-2002;
29-MAY-2002;
06-JUI-2002;
06-JUI-2002;
07-AUG-2002;
20-AUG-2002;
23-AUG-2002;
23-AUG-2002;
23-AUG-2002;
23-AUG-2002;
24-AUG-2002;
26-AUG-2002;
                                                                                                       Agee ML, Alsobrook JP, Anderson DW, Berghs C, Boldog FL; Burgess CE, Catterton E, Dipippo VA, Edinger SR, Bisen A; Ellerman K, Gangolli EA, Gerlach V, Gorman L, Rothberg BG, Gi Herrmann JL, Halvorsen Y, Ji W, Kekuda R, Khramtsov NV; Larochelle WJ, Lepley DM, Li L, Macdougalli JR, Miller CE, Ort Padigaru M, Patturajan M, Pena CEA, Peyman JA, Rieger DK; Rothenberg ME, Shenoy SG, Smithson G, Spaderna SK, Spytek KA; Stone DJ, Taupier RJ, Vernet CAM, Voss EZ, Zhong M;
                                                                 WPI;
New human NOVX polypeptides and nucleic acid molecules, useful diagnosing, preventing or treating NOVX-associated disorder, e.
                                                                                                                                                                                                                                                                                                                                                                         (AGEE)
(AAGE)
(AAGE)
(ANDE)
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(BERG)
(BOLDY)
(ELISE)
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                                                                      2004-268786/25
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SHENOY S G.
SMITHSON G.
SMITHSON S K.
SPADERNA S K.
SPYTEK K A.
STONE D J.
TAUPIER R J.
TRUPIER R J.
VERNET C A M.
VOSS E Z.
ZHONG M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PADIGARU M.
PATTURAJAN M
PENA C E A.
PEYMAN J A.
RIEGER D K.
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BLLERMAN K.
GANGOLLI B J
GERLACH V.
GORMAN L.
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BERGHS C.
BOLDOG F L.
BURGESS C E.
CATTERTON E.
DIPIPPO V A.
EDINGER S R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GUO X S.
HERRMANN J
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ALSOBROOK J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KHRAMTSOV N V.
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2002US-0381004P
2002US-0381495P
2002US-0383534P
2002US-0383744P
2002US-0383829P
2002US-0383829P
2002US-0393332P
2002US-0401315P
2002US-0401315P
2002US-0401788P
2002US-0404670P
2002US-040569P
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2002US-040569P
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2002US-040569P
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e.g. cancer, for

Ort

Guo Ή. SX

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RESULT 770
AAQ67183/c
ID AAQ67183;

XX
AC AAQ67183;

XX
DT 25-MAR-2003 (revised)
DT 22-MAR-1995 (first entry)
XX
DE Primer for amplifying part of th
XX
Immunisation; vaccine; therapy;
KW non-functional gene; template; a
KW human immunodeficiency virus; ss
XX
OS Synthetic.
XX
WO9416737-A1.
XX
PD 04-AUG-1994; 94WO-US000899.
XX
PP 26-JAN-1994; 94WO-US000899.
XX
PP 26-JAN-1993; 93US-00008342.
PR 11-MAR-1993; 93US-00008343.
PR 11-MAR-1993; 93US-00029336.
PR 11-MAR-1993; 93US-00029336.
PR 21-SEP-1993; 93US-00029335.
PR 21-SEP-1993; 93US-00124962.
PR 21-SEP-1993; 93US-00125012.
XX
PA (WEIN/) WEINER D B.
PA (WAIN/) WILLIAMS W V.
PA (WANG B.
PA (CONE/) CONEY L R.
PA (MANG/) MANG B.
PA (CONE/) ZURAWSKI V R.
XX
DR Weiner DB, Williams WV, Wang E
XX
DR WPI; 1994-263787/32.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to human NOVX polypeptides and the polynucleotides CC encoding them. The invention also relates to antibodies specific to the concoding them. The polypeptides, polynucleotides and antibodies are CC NOVX polypeptides. The polypeptides, polynucleotides and antibodies are useful for manufacturing a medicament for treating a syndrome associated with a human disease, such as a pathology associated with the NOVX CC polypeptide. The sequences are useful for diagnosing, treating or CC preventing a NOVX-associated disorder, e.g., cancer, atherosclerosis, CC disease, scleroderma, hypertension, haemophilia, idiopathic consist, CC disease, scleroderma, hypertension, haemophilia, idiopathic CC chrombocytopenic purpura, immunodeficiencies, AIDS, dyslipidemia, CC cachexia, multiple sclerosis or fertility. The nucleic acids may be used CC cachexia, multiple sclerosis or fertility. The nucleic acids may be used CC cachexia, multiple sclerosis or fertility. The sclerosis as the properties of the properties of pharmacogenomics. This sequence represents a CC preventive medicine or in pharmacogenomics. This sequence represents a CC preventive medicine or in pharmacogenomics.
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 5 A; 4 C; 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunisation; vaccine; therapy; prophylaxis; defective gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  scleroderma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
les 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4596 ACTGCATGGACAGGTGC 4612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amplifying part of the gag region of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
ilarity 94.1%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        gene; template; antisense; ribozyme; bupivacaine; HIV;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NO 382;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       virus; ss
                                                                  Wang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      610pp;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.4;
Pred. No. 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 8.70
); Mismatches
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                                                                     Coney LR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0 Other;
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                                                                       Merva MJ,
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                                                                     Zurawski
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FFFX8X55555555555555555555555
                                                                                                                                                  Method for introducing genetic material into cells - utilises polynucleotide function enhancer and nucleic acid free of retroviral particles, e.g. HIV immunisation.
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Page 109; 136pp; English.

(bupivacaine) and (b) administering to the cells the nucleic acid molecule free of retroviral particles. Nucleic acid molecules which are delivered to cells may serve as genetic templates for proteins that function as prophylactic and/or therapeutic immunising agents; replacement copies of defective, missing or non-functional genes; genetic templates for therapeutic proteins; genetic templates for antisense molecules or as genetic templates for ribozymes. Two primers (AAQ67182, AAQ67183) were used to amplify part of the gag coding region from a plasmid construct which comprised the 5' LTR and the rest of the HIV-1 genome to nucleotide 5795 (Genbamk numbering) cloned into the XBaI and SalI sites of Bluescript (The HIV-1 sequences are obtained from the HXB2D plasmid). The amplified sequence was used in the construct designated pGAGPOL rev2, used to express HIV gag and pol genes. (Updated on 25-MAR-2003 to correct PN field.) Genetic material may be introduced into the cells of an individual by contacting the individual's cells with a polynucleotide function enhan al by (a)

Sequence 21 BP; 0 A; 4 C; 10 G; 7 T; 0 U; 0 Other;

Query Match Best Local : Matches l Similarity
16; Conserv Conservative 94.1%; 0.3%; 0; Score 15.4; Pred. No. 8 Mismatches 8.8e+02 몂 1; Length 21; Indels 0 Gaps 0

맑 á 1261 AGCCTACAGCCCCACCA 1277 N

AAT00869 standard; DNA; 21 ВP

16-MAY-1996 (first entry)

HIV strain HXB2 gag coding region 3' primer.

RESULT 771
AATOO869/c
ID AATOO8
XX AATOO8
XX IMMUNI
XX IMMUNI
XW INTERIOR
XX IMMUNI
XW INTERIOR
XX IMMUNI
XW INTERIOR
XX IMMUNI
CCC INTERIOR
CCC Immunisation; disease; pathogen; genetic vaccine facilitator; saponin; anionic lipid; lectin; oestrogen; alkyl; dimethylsulphoxide; urea; PCR; retroviral particle; retrovirus; HIV; SIV; epitope; primer; amplification; 88

Synthetic

WO9526718-A1

12-OCT-1995

30-MAR-1995; 95WO-US004071.

01-APR-1994; 94US-00221579

(APOL-) APOLLON INC

₽,

WPI; 1995-358434/46.

Introducing genetic material into cells of an individual -the cells with a genetic vaccine facilitator and a nucleic by contacting acid molecule

19, Page 50; 114pp; English.

Immunisation of an individual against a disease or pathogen comprises introducing genetic material (a genetic vaccine) into the cell of the individual by contacting the cell with a genetic vaccine facilitator

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RESULT 772
ACT ANTI3768
ID ANTI3767
AC ANTI37
AC (UYDU-PA (HSCR-XX)
AC (UYDU-PA (HSCR-XX)
AC (UYDU-PA (HSCR-XX)
AC ANTI3
AC ANTI3
CC ANTI3
CC ANTI3
CC ANTI3
CC ANTI3
CC ANTI3
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SQ Seque
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Matches
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Best Local Similarity
Matches 16; Conserv
                    Query Match
Best Local Similarity
                                                                                                        Oligonucleotides derived from introns of the endoglin gene can be used primers for amplifying a single exon of the endoglin gene for its use i diagnosis of haemorrhagic telanglectasia (HHT). DNA encoding endoglin c be used for gene therapy of HHT which is caused by inheritance of a defective gene, e.g. endoglin, beta-glycan, TGP-beta type I or II receptor or TGF-beta/activin type I receptor. Two primers (AAT13767, AAT13768) were used to amplify an exon sequence of the endoglin gene. Primer pairs used in this method are described in AAT13757-78
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (GVF) selected from anionic lipids, saponins, lectins, oestrogenic cpds., hydroxylated lower alkyls, dimethylsulphoxide (DMSO) or urea, and a nucleic acid that is free of retroviral particles. The primers AAT00830-71 are used in the construction of the genetic vaccines based on sequences of HIV, SIV or pathogenic bacterial epitopes. The primers AAT00868-9 were used to amplify part of the gag coding region from HIV strain HXB2 for construction of the genetic vaccine plasmid pGACPOL rev2. The plasmid contains the rev gene from HIV strain HXBB and the gag open reading frame (ORF), part of the pol ORF and the rev response element from HIV strain
                                                                                                                                                                                                                                                                                                                              Human gene for endoglin (t useful in diagnosis and
                                                                                                                                                                                                                                                                         Disclosure; Page 13; 71pp; English.
                                                                                                                                                                                                                                                                                                              telangiectasia.
                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1996-286827/29.
                                                                                                                                                                                                                                                                                                                                                                                                                          Marchuk DA, Mcallister K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         06-JUN-1996.
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gene therapy; haemorrhagic telangiectasia; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Primer for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-DEC-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16;
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   16;
                                                                         21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      amplifying endoglin exon sequence.
                                                                      BP; 6 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
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   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                 94.1%;
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                                   0.3%;
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                                                                                                                                                                                                                                                                                                                              (transforming growth factor beta binding protein)
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Score 15.4; DB 1
Pred. No. 8.8e+02
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.4; DB 1;
Pred. No. 8.8e+02;
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                                   DB 1; Length 21;
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RESULT 773
AAZ18406/c
ID AAZ184
XX AAZ184
XX AAZ184
XX POlymo
XX POlymo
XX ASTH1;
XW CHERAP
XX HOMO 6
XX HOMO 6
XX HOMO 7
XX HOMO 7
XX WO9937
XX WO9937
XX WO9937
XX ELJAN
XX AAXTH1;
CC MARTMAL
PF WAMMMAL
PF WAMMAL
PF WAMMMAL
PF WAMMAL
PF WAMMMAL
PF WAMMAL
PF WAMMMAL
PF WAMMAL
PF WAMMMAL
PF WAMMMA
RESULT 774
AAZ97503
ID AAZ975
XX
AC AAZ975
XX
AC AAZ975
XX
DT 15-SEP
DT 26-APR
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Best Local
                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention identifies a genetic locus ASTH1, associated with asthma, mapped to human chromosome 11p. ASTH11 and ASTH11 are genes present within the locus, located close to each other on human chromosome 11p, and have similar patterns of expression, and common sequence motifs. The ASTH1 genes and fragments, encoded protein, genomic regulatory regions and anti-ASTH1 antibodies are useful in the identification of individuals predisposed to development of asthma, and for the modulation of gene activity in vivo for prophylactic and therapeutic purposes. The ASTH1 protein is useful as an immunogen to raise specific antibodies, in drug screening for compositions that mimic or modulate ASTH1 activity or expression, including altered forms of ASTH1 protein, and as a therapeutic. Sequences AAZ18366-Z18509 represent polymorphisms in the
  15-SEP-2003
26-APR-2000
                                                                                 AAZ97503;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                  AAZ97503 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 62; 195pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mammalian asthma related genes, useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polymorphic fragment in region 5' to ASTH1J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ASTH1; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    development of asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2609
                                                                                                                                                                                                                                                                                                                   4891 ACAAGTTGCCATCTGGTTT 4909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1999-479058/40.
                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and ASTH1J genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ω
                                                                                                                                                                                                                                                                                                                                                                                               Similarity
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(revised)
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immunogen; polymorphism;
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                                                                                                                                     DNA;
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                                                                                                                                     21
                                                                                                                                     ВP
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                                                                                                                                                                                                                                                                                                                                                                                                  Score 15.4;
Pred. No. 8
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                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                    .8e+02;
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AAZ975
AAZ975
ID AAZ975
XX AZ975
XX 15-SEP
DT 15-SEP
DT 26-APR
XX DB HIV-1
XX drug 8
KW drug 8
XX drug 9
XX drug 10
XX drug 10
XX human
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention describes the detection of drug-selected mutations in the HIV protease gene. The method of detection allows the simultaneous characterisation of a range of codons involved in drug resistance using sets of probes optimised to function together in a reverse-hybridisation assay. AAZ97517 to AAZ97997 represent specifically claimed probes for use in the assay, and AAZ97479 to AAZ97501 represent specifically claimed HIV protease gene polymorphic nucleotide sequences. AAZ97502 to AAZ97515, and AAZ98004 to AAZ98007, represent PCR primers for the HIV protease gene, and AAZ97516 represents an HIV protease probe used in an example from the present invention. The method, probes and primers can be used for the detection of drug-selected mutations in the HIV protease gene. The method allows the simultaneous characterisation of a range of codons involved in drug resistance. The method may also be used for HIV protease genotyping assays. The probes are able to discriminate between wild type and mutated protease sequences. The method allows rapid and reliable detection of drug-selected mutation in HIV. (Updated on 15-SEP-2003 to standardise OS
                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
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                                                   Human immunodeficiency virus; HIV; prot drug selected mutation; hybridisation;
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   Human immunodeficiency virus 1.
                                                                                                     HIV-1 protease gene PCR primer SEQ ID NO:502.
                                                                                                                                      15-SEP-2003
26-APR-2000
                                                                                                                                                                                          AAZ97507;
                                                                                                                                                                                                                        AAZ97507 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HIV-1 protease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       n immunodeficiency virus; HIV; protease; probe; detection; selected mutation; hybridisation; genotyping; infection; resistance; PCR primer; ss.
                                 resistance; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             4; Page 19; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                               16;
                                                                                                                                                                                                                                                                                                                                                                                               n 0.3%;
Similarity 94.1%;
                                                                                                                                                                                                                                                                                                                                                                               Conservative
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(first entry)
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                                                                                                                                                                                                                        DNA;
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                                                 protease; probe; detection; ion; genotyping; infection;
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The present invention describes the detection of drug-selected mutations CC in the HIV protease gene. The method of detection allows the simultaneous CC characterisation of a range of codons involved in drug resistance using CC sets of probes optimised to function together in a reverse-hybridisation CC assay. AAZ37917 to AAZ37997 represent specifically claimed probes for use CC in the assay, and AAZ37997 represent specifically claimed probes for use CC protease gene polymorphic nucleotide sequences. AAZ37902 to AAZ39004 to AAZ38007, represent PCR primers for the HIV protease gene, CC and AAZ37516 represents an HIV protease probe used in an example from the CC present invention. The method, probes and primers can be used for the CC allows the simultaneous characterisation of a range of codons involved in CC drug resistance. The method may also be used for HIV protease genotyping CC assays. The probes are able to discriminate between wild type and mutated CC protease sequences. The method allows rapid and reliable detection of CC drug-selected mutation in HIV. (Updated on 15-SEP-2003 to standardise OS field)
Sequence 21 BP; 7 A; 10 C; 4 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Page 20; 76pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of drug-selected mutations in the HIV protease gene used treat HIV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-147219/13
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Matches Query Match Best Local (16; Conserv Conservative 0.3%; 94.1%; 0 Score 15.4; DB 1 Pred. No. 8.8e+02 Mismatches DB 1; Length 21; Indels 0, Gaps 0

밁 Ś 1261 AGCCTACAGCCCCACCA 1277 4 20

10-SEP-2001 AAZ77203 standard, DNA; 21 (first entry) ВP

Human biallelic marker downstream amplification primer SEQ ID NO:11559.

RESULT 776
AAZ77203/c
ID AAZ772
XX
AC AAZ772
XX
Human
DB Human
XX
Human
KW Human
KW Haplott
KW haplott
KW diagno
XX
Human
XX
W Haplott
XX
Human
XX Human genome; biallelic marker; high density disequilibrium map; genomic map; haplotype; phenotype; polymorphic base; genotyping; haplotyping; hybridisation; identification; characterisation; amplification; single nucleotide polymorphism; SNP, PCR nrimar. 21-APR-1999; 28-OCT-1999. diagnosis; ss Homo sapiens 99WO-IB000822

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RESULT 777
AAZ95599
ID AAZ9559
XX AAZ955
XX AC AAZ955
XX Human
XX Human,
XW Human,
XW Osler-
XX Homo &
XX US6022
XX 
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Matches 16
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N.B. The SEQ ID NOS 2852, 2913, 2974, 3035, 3096, 3157, 3227, 3297 and 3367, are not accually given a sequence in the Sequence Listing from the present invention.
                            Diagnosing hereditary hemorrhagic telangiectasia by identifying genetic mutations that lead to susceptibility to the disease.
                                                                                                                                                                                                                                                                                                                                                                                                08-FEB-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       US6022687-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; endoglin; hereditary haemorrhagic telangiectasia; HHT;
Osler-Weber-Rendu disease; diagnosis; identification; PCR pri
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human endoglin PCR primer SEQ ID NO:31.
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23-NOV-1998;
                                                                                                                                                                    Marchuk DA,
                                                                                                                                                                                                                                                                                  29-NOV-1994;
                                                                                                                                                                                                                                                                                                                                      29-NOV-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens
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                                                                                                                                                                    Mcallister K,
                                                                                                                                                                                                                           DUKE
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98US-0109732P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15.4; DB 1. Pred. No. 8.8e+02
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Matches 16
             consisting of a cis-acting mRNA transport sequence (RTS) (Ia) of a myelin basic protein (MBP) cDNA or its portion that enhances transport of mRNA within the cytoplasm, or consisting of a cis-acting mRNA localisation enhancer (RLE) (Ib) of MBP cDNA or its portion which enhances localisation of mRNA within the cytoplasm. (I) placed 5' or 3' to a leterologous gene that is operably linked to a 5' regulatory region and a 3' termination sequence, in an expression vector is useful for expressing a heterologous gene which involves transforming a host cell with the expression vector. (I) confers properties such as localisation, transport and increased translation efficiency of a heterologous mRNA transgript when transcribed into such mRNA. (I) present in a vector is useful for increasing translation of a heterologous gene, by which increased
                                                                                                                                                                                                                                                                                                              Novel nucleic acid molecule having cis-acting mRNA transport sequence localization enhancer which enhance transport or localization of mRNA within cytoplasm, respectively, is useful for enhancing gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a method (I) for diagnosing, or identifying a predisposition to, hereditary haemorrhagic telangiectasia (HHT) (also called Osler-Weber-Rendu disease) comprising determining whether a sample of genomic DNA from the subject contains a mutation in the gene encoding endoglin (the mutation is indicative of a predisposition to HHT). The method may be used for diagnosing (or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Myelin basic protein; MBP; cis-acting mRNA transport sequence; RTS; cis-acting mRNA localisation enhancer; RLE; gene therapy; localisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 6 A; 4 C; 8 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             predisposition to HHT). The method may be used for diagnosing (or identifying individuals with a predisposition to developing) HHT. The present sequence represents a PCR primer for the human endoglin gene,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Col 7-8; 40pp;
                                                                                                                                                                                                                                         The present invention describes an isolated nucleic acid molecule
                                                                                                                                                                                                                                                                                                                                                                                                                          Carson J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US6225082-BI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transport; translation; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bovine RTS homology transported mRNA NOS SEQ ID NO:11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH21622;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAH21622 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               which is used in the exemplification of the present invention
increasing translation of a heterologous production of recombinantly produced prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                              (RESE ) RESEARCH CORP TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2609 AGGGGAGGAACCTGATG
                                                                                                                                                                                                                                                                                                                                                                                        2001-307679/32.
                                                                                                                                                                                                                                                                           3; Fig 3; 39pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AGGGGAGGAACCAGATG
                                                                                                                                                                                                                                                                                                                                                                                                                        Kwon S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                          Ainger K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
94.1%;
                                                                                                                                                                                                                                                                             English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 15.4;
Pred. No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                            Avossa D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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នុក្ស
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Best Local S
Matches 16
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Guanidinium functionalized oligomers prepared from corresponding monome: units, are hybridizable with a specific RNA or DNA sequence, useful for diagnostic and therapeutic purposes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     research scale or on a large commercial scale is increased. The recombinantly produced proteins whose translation is enhanced by (I) may be a biologically active protein, structural or therapeutic protein. (I) is useful in gene therapy. The present sequence represents a transported mRNA sequence with RTS homology, which is used in the exemplification of
        AAF96199;
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 21 BP; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present invention relates to nucleotide oligomers comprising monomounits. Oligomers modulate gene expression when hybridized by a single-double-stranded nucleic acid. They are useful for gene therapy,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          07-JUL-2000; 2000WO-US018609
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                                                        AAF96199 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                            diagnostic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 36; Page 62; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-138119/14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Manoharan M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           11-JAN-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gene expression; gene therapy; diagnosis; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-APR-2001
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                                                                                                                                                                                     AGAGAGAGAAAAATC 2
                                                                                                                                                                                                                                                                                                                                                                                                                                            and investigative purposes
                                                                                                                                                                                                                                                                                   0.3%;
llarity 94.1%;
Conservative
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llarity 94.1%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                         A; 7 C; 2
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                                                                                                                                                                                                                                                                                                                                                                                         G; 10 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                      Score 15.4; DB 1;
Pred. No. 8.8e+02;
0; Mismatches 1
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0; Mismatches
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Pred. No. 8.
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RESULT 781
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                               The present invention provides a method of diagnosing a vascular disease in an individual, involving determining the sequence at various polymorphic sites within the human thrombospondin 1 and thrombospondin 4 genes. The sequences at a number of polymorphic sites are also provided in the specification. In particular, the method can be used in the disease, stroke, peripheral vascular diseases, venous thrombosembolism and pulmonary embolism. Single nucleotide polymorphisms (SNPs) are also useful in forensics, paternity testing, genetic analysis and phenotype correlations to diseases. The present sequence is an example of one of the human gene SNPS shown in the specification
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleic acids comprising single nucleotide polymorphisms, useful applications such as forensics, paternity testing, medicine, generallysis and phenotype correlations to diseases such as diabetes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        10-SEP-1999; 99US-0153357P.
26-JUL-2000; 2000US-0220947P.
16-AUG-2000; 2000US-0225724P.
                            24-OCT-2001
                                                                                AAS11636 standard;
                                                                                                                                                                                                                                                        Sequence 21
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                                                                                                                                                                                                                  Similarity
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                                                                                                                                                AGAGAGGGCGGATTCCC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             single nucleotide
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                                                                                                                                                                                                      Conservative
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                          (first entry)
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/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /standard_name= "single nucleotide polymorphism"
                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           paternity test; ds.
                                                                                                                                                                                                                  94.1%;
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Pred. No. 8.8e+02
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Human CYP2B6 allele sequencing primer seqCYP2B6-P2F for promoter region.

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ARESULT 782
AAP27878/C
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        predisposition to several common cancers caused by environmental carcinogens, and diseases treated with drugs that are targets of the CAPCER6 gene product, whose metabolism is therefore dependent on CYP2B6. Cancer or susceptibility to cancer can be diagnosed by detecting the presence of a molecular variant of CYP2B6. From variants of the alleles, modulators of the activity can be developed for use in treatment and prevention of CYP2B6-related disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The sequence represents a sequencing primer used to sequence the promoter region of the human CYP2B6 gene. It is used for specific detection and genotyping of CYP2B6 alleles in humans, determination of which is useful for the optimisation of therapies utilising CYP2B6 substrates. Oligonucleotide sequences are useful in detection of the individual
                                                                                                                                                                                                                                                                             cerebroprotective; immunomodulator; vulnerary; vasotropic; gene therapy; hyperplasia; tumour; restenosis; psoriasis; Dupuytren's contracture; diabetes; rheumatoid arthritis; cerebral oedema; Alzheimer's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF27878 standard; DNA; 21
      01-JUN-2000; 2000WO-US015303
                                                                  14-DEC-2000
                                                                                                                        WO200075321-A2
                                                                                                                                                                                                                                                                                                                                                                  Human; NOVX; antiinflammatory; cytostatic; neuroprotective;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Human NOV7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-MAR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF27878;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New polynucleotide(s) of the polymorphic human CYP2B6 gene for detection and treatment of disorders i.e. cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      09-FEB-2000; 2000EP-00102701
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            09-FEB-2001; 2001WO-EP001456
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200159152-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EPID-) EPIDAUROS BIOTECHNOLOGIE
                                                                                                                                                                                                                                             primer; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4869 GTCTCAGTTTCTTCTC 4885
                                                                                                                                                                                       sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lang T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 1 A; 7 C; 3 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                 forward primer
                                                                                                                                                                                                                                                88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.4; DB 1; Length 21; Pred. No. 8.8e+02;
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ACD40283/c
ID ACD402
XX ACD402
AC ACD402
XX Tissue
XW Pancre
XW Cardic
XW Lische
XW Huntri
XW Huntri
XW ALS; F
XW Gene t
XX
PN US200:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence was used to isolate a new isolated polypeptide (NOVX). The NOVX polypeptides, NOVX nucleic acids, and anti-NOVX cc antibodies are useful for treating or preventing NOVX-associated cd disorders. They are also useful for determining the presence of or a predisposition to a disease associated with altered levels of the NOVX copypeptide or nucleic acid. These NOVX-associated disorders include hyperplasias, tumours, restenosis, psoriasis, Dupuytren's contracture, cc diabetic complications, rheumatoid arthritis, cerebral lesions, diabetic neuropathies, cerebral oedema, senile dementia or Alzheimer's disease. Cc The NOVX polynucleotides are especially useful in gene therapy. Specifically, NOVX is useful as a diagnostic marker or prognostic marker, protein therapeutic and antibody target or small molecule drug target to treat disorders in the immune response pathway, thyroid and metabolic disorders, diseases of the pancreas (e.g. cdiabetes or digestive disorders), proliferative diseases, or tissue regeneration and development (e.g. wound healing or treatment of burns) or complete the complete or segmentation and development (e.g. wound healing or treatment of burns) or regeneration and development (e.g. wound healing or treatment of burns)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 16; Conserv
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16-MAR-2000; 2000US-0189810P.
22-MAR-2000; 2000US-0191158P.
30-MAR-2000; 2000US-0193086F.
31-MAY-2000; 2000US-00137322.
                                                                                                                                                                                                                                                                                                        Tissue typing; cancer; breast cancer; colon cancer; lung cancer; sarcoma; pancreatic cancer; uterine cancer; organ transplantation disorder; ss;
                                                                                          ALS; peripheral neuropathy; nervous system tumour; neurotoxin; tremor; neuropathy; acute brain injury; peripheral nerve trauma; human; NOVX; gene therapy; epilepsy; breast tumour associated protein 47.
                                                                                                                                                            Cardiovascular disease; melanoma; atherosclerosis; diabetes mellitus; ischaemic heart disease; haemorrhage; peripheral vascular disease; PCR; ithrombosis; hypertension; systemic lupus erythematosus; haematopoiesis; tissue regeneration; wound healing; hyperproliferative disorder; primer; psoriasis; neural disorder; Parkinson's disease; Alzheimer's disease; Buntington's disease; multiple sclerosis; amyotrophic lateral sclerosis;
                                                                                                                                                                                                                                                                                                                                                                          Breast tumour associated protein 47-like polypeptide NOV7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ACD40283 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 8 A; 4 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New NOVX polypeptides and polynucleotides, useful in gene therapy, as a diagnostic marker, protein therapeutic, antibody or small molecule drug target for treating immune, proliferative and metabolic diseases and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (CURA-) CURAGEN CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.4; DB 1;
Pred. No. 8.8e+02;
0; Mismatches 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21;
                                                                                                                                                                                                                                                                                                                                                                                 forward primer.
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RESULT 784
ADH97217/c
ID ADH972
XX
AC ADH972
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                                                                                                                                                                                                                                                                                                                                                    Creatment including therapeutic and prophylactic. The nucleic acid is also useful for expressing NOVX protein. The nucleic acid is also useful for expressing NOVX protein. The nucleic acid is also useful to provide polynucleotide reagents e.g. labelled probes that are useful in an in situ hybridisation technique, for identifying a specific tissue carid is also useful for many thing energy on a chromosome and thus locating gene regions associated with genetic disease, identifying an an individual carid is also useful for mapping genes on a chromosome and thus locating gene regions associated with genetic disease, identifying an individual carid is also useful for treating cancer. The nucleic acid is also useful for treating cancer, ce specially cancers of the breast, colon, lung, pancreas or uterus, or a chromosome or sarcoma. The nucleic acid is also useful for treating cancer, ce atherosclerosis, ischaemic heart disease, hamenorphage, diabetes mellitus, peripheral vascular disease, thrombosis, hypertension and systemic lupus cregulating haematopoiesis, for regeneration of bone, cartilage, tendon cartilage, The nucleic acid is also useful for treating cancer. The nucleic acid is also useful for treating cancer, complained and concern encoded by the nucleic acid is useful for cregulating haematopoiesis, for regeneration of bone, cartilage, tendon cartilage, tendon cancer and/or nerve tissue growth or regeneration and for wound cancer and/or nerve tissue growth or regeneration and for wound cancer and and systemic and systemic cancer and concerns the nerve at a secure and the neuropathies, peripheral nerve trauma or injury, and other neuropathies, cancer and other neuropathies, cancer and concerns and/or tremors. The present sequence represents a human breast cumour associated protein 47-like polypeptide PCR primer
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Best Local S
Matches 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New nucleic acid sequence encoding a human breast tumor-associated protein 47-like polypeptide, useful for treating cardiovascular disorders, neural disorders, diabetes mellitus and cancers.
                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 8 A; 4 C; 6 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 22-MAR-2000; 2000US-0191158P.
30-MAR-2000; 2000US-019306BP.
03-MAY-2000; 2000US-0201188P.
31-MAY-2000; 2000US-00584411.
                      ADH97217
                                                                 ADH97217 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a new isolated NOV4 nucleic acid. The nucleic
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22-MAR-2000;
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                                                                                                                                                                                                                     1925 CTTCTTTGGAGCAGGCA 1941
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                                                                                                                                                                                                                                                                                       Similarity
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                                                                                                                                                                                                                                                                                  94.1%;
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                                                                                                                                                                                                                                                                                                         Score 15.4;
                                                                                                                                                                                                                                                                                       Pred.
                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                     No. 8.8e+02
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                                                                                                                                                                                                                                                                                                         DB 1; Length 21;
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ADJ13812/c
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention relates to novel synthetically modified oligomers that have increased nuclease resistance and have enhanced hybrid binding. Such oligomers are useful for diagnostic and therapeutic uses such as antisense technologies. The invention also discloses a method for the preparation of the oligomers with modifications as fully defined in the specification. The present sequence represents a synthetically modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nuclease resistance; hybrid binding; antisense technology; ss
                                                                                                                       27-JUN-2001; 2001US-0301370P
                                                                                                                                                                             27-JUN-2002; 2002US-00184085
                                                                                                                                                                                                                                         14-AUG-2003
                                                                                                                                                                                                                                                                                               US2003152950-A1
                                                                                                                                                                                                                                                                                                                                                                                                          tumour suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human DNA probe used to immobilise CpG methylated DNA SeqID 939.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADJ13812;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21 BP; 2 A; 7 C; 2 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        specification. The present sequence represents a synthetically modified oligonucleotide of the invention.
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                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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                                   (GARN/) GARNER H R. (MINN/) MINNA J D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1186 AGAGAGAGAGAGAAATC 1202
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                                                                                                                                                                                                                                                                                                                                                                                                                                        ss; chemical modification; methylation; array; CpG island
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ilarity 94.1%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.4; DB 1;
Pred. No. 8.8e+02;
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RESULT 786
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XX ADJ131
XX DT 20-MAY
XX Probe;
XW Elmon a
XX Elmon a
XX US2003
XX GARN,
PA (LUEB,
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Matches 16
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 Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
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                                                                    WPI; 2003-874843/81.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Human DNA
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                                                                                                                                      (BALO/)
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                                                                                                                                  GARNER H R.
MINNA J D.
LUEBKE K J.
BALOG R P.
                                                                                                                                                                                                                                                                                                                                                                                       suppressor;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      probe used to immobilise CpG methylated DNA SeqID 233.
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                                                                                                     Luebke
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No. 8.8e+02;
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RESULT 787
ADJ13699/C
ID ADJ136
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XX YX ADJ136
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DE Human
XX YX Lumour
XX YX ADJ136
PF 27-JUN
XX 27-JUN
XX 27-JUN
XX 27-JUN
XX ADJ196
PF 27-JUN
XX ADJ197
PA (MINN)
PA (LUEB)
PA (LUEB)
PR (DAILO)
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               This invention relates to a novel method for analysing chemically modified macromolecules. Specifically, it refers to a high throughput method for the parallel analysis of many potential sites of chemical modification (e.g. methylation) in DNA. The present invention describes treating the DNA with one or more chemical reagents that result in
                                                                                                                                                               Analysis of chemical modification of DNA involves obtaining sample of DNA to be analyzed, treating DNA with chemical reagents that result in different base sequences, and determining sequence of resulting DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human
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                                                                                                                                                                                                                                                                          Garner HR,
                                                                                                                                                                                                                                                                                                                                                  (GARN/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                           27-JUN-2002; 2002US-00184085.
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                                                                                                                                                                                                                                                                                                                                   (LUEB/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA
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) MINNA J D.
) LUEBKE K J.
) BALOG R P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        suppressor; p16; human; H69; H1618.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ss; chemical modification; methylation; array; CpG island
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                                                                                                                           1; SEQ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             probe used to immobilise
                                                                                                                                                                                                                                                                            Minna JD,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 4 A;
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                                                                                                                                                             l, treating D
sequences,
                                                                                                                             ID NO 826;
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                                                                                                                                                                                                                                                                            Luebke
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                                                                                                                                                                                                                                                                              RP;
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16-MAR-2000;
22-MAR-2000;
The invention relates to an isolated nucleic acid encoding a human cell adhesion molecule-like protein, comprising a sequence encoding a polypeptide having a sequence appearing as ADM55387, a sequence at least 90% identical to the nucleic acid, a sequence encoding a polypeptide having conservative amino acid substitutions to the protein or a fragment
                                                                                                                                                                                                                       New nucleic acids encoding human cell adhesion molecule-like proteins, useful for treating e.g. cancers, neurological disorders, viral, bacterial, fungal, helminthic and protozoal infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          allergic reaction; autoimmune haemolytic anaemia; rheumatoid viral infection; bacterial infection; fungal infection; helminthic infection; protozoal infections; primer; RTQ-PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    lymphoma; melanoma; neurological disorder; epilepsy; ischaemic cerebrovascular disease; Alzheimer's disease; ischaemic cerebrovascular disease; Alzheimer's disease; pick's disease; pestic fibrosis; pick's disease; golter; gastrointestinal disorder; diabetes mellitus; Grave's disease; golter; gastrointestinal disorder; dicerative colitis; gastric ulcer; duodenal disorder; autoimmune disease; ulcerative colitis; gastric ulcer; duodenal disorder; autoimmune disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modification of interest. Accordingly, a device comprising an array of probes is provided to hybridise with and select the altered DNA sequences that comprise the modifications of interest such as a CpG island. In particular, this invention refers to analysing the methylation pattern of a region of the promoter for the tumour suppressor gene pl6 from two human lung tumour cell lines H69 and H1618. This oligonucleotide sequence is a human DNA probe used to immobilise CpG methylated DNA of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human cell
                                                                                                                                                                         Example
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; ss; PCR; cell adhesion molecule; NOVX; cancer; leukaemia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 21 BP; 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                              (CURA-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURAGEN CORP.
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                                                                                                                                                                       SEQ
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2000US-0189810P.
2000US-0191158P.
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2000US-0201388P.
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                                                                                                                                                                    78pp;
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Pred. No. 8.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            G; 5
                                                                                                                                                                    English.
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                                                              xx antiarthritic; antiatheroscierotic; haemostatic; vasotropic; xx thrombolytic; antidiabetic; hypotensive; dermatalogical; xx thrombolytic; antidiabetic; hypotensive; dermatalogical; xx immunosuppressive; antiinflammatory; immunostimulant; fungicide; xx virucide; protozoacide; neuroprotective; antirheumatic; antiarthritic; xx antiasthmatic; antiparkinsonian; nootropic; anticonvulsant; xx virucide; protozoacide; neuroprotective; antirheumatic; antiarthritic; xx antiasthmatic; antiparkinsonian; nootropic; anticonvulsant; xx virucide; yeartrophy; osteoarthritis; xx virucidessis; keloid; xx virucidessis; keloid; xx virucidessis; keloid; xx virucidessis; keloid; xx virucidessis; papertrophy; osteoarthritis; xx virucidessis; diabetes mellitus; hypertransion; hypothyroidism; xx virucidessis; autoimmune disorder; connective tissue disease; xx virucidessis; systemic lupus, erythematosus; rheumatoid arthritis; xx virucidessis; systemic lupus, erythematosus; rheumatoid arthritis; xx virucidessis; insulin dependent diabetes mellitus; xx virucidessis; yraft- versus-host disease; xx haematopoiesis; tissue regeneration; vound healing; tissue repair; burn; incision; ulcer; xx virucidesse; xx vi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Huntington's diseas human; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADF66780 standard; DNA; 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local
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16; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hepatotropic; vulnerary; antipsoriatic; osteopathic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein NOV7
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                                              disease; amyotrophic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   53
                                              Parkinson's disease;
sclerosis; Shy-Drager
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 21;
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                                                                                                                             incision; ulcer;
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                                              syndrome;
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RESULT 790
ADF91944/c
ID ADF919
XX
AC ADF919
XX
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                                                                                                                                                                                                                        Query Match
Best Local S
Matches 16
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30-MAR-2000;
03-MAY-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention describes a substantially purified polypeptide (I) having amino acid sequence chosen from a fully defined NOVX sequence (S1) of 708 amino acids as given in the specification, or polypeptide having one or more conservative amino acid substitutions of (S1), or mutant or variant of (S1). (I) having (S1) is useful for diagnosing a pathological condition associated with (I) or its activity in a subject e.g. cancer. (I) useful in treatment of cancer, hyperproliferative diseases,
                   ADF91944;
                                                                                                                                                                                                                                                                                                                                amyotrophic lateral sclerosis, and Shy-Drager syndrome. (I) has effectiv
antitumour and antiinflammatory activity. This sequence represents a
primer used in the isolation of DNA encoding a novel human NOVX protein.
                                                       ADF91944 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                 Sequence 21 BP;
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; 2000US-01198110P.
; 2000US-0119158P.
; 2000US-0193086P.
; 2000US-0201388P.
; 2000US-005844111.
                                                                                                                                                                                                                            Conservative
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                                                                                                                                                                                                                                           94.18;
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                                                                                                                                                                                                                                                            Score 15.4;
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                                                                                                                                                                                                                                                           Length 21;
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RESULT 791
ADG47997/c
ID ADG47997 standard; DNA; 21 BP
XX
AC ADG47997;
XX
DT 11-MAR-2004 (first entry)
XX
DE Duplex DNA strand 2 used for
XX
DE Upplex DNA strand; the
XX
OS Unidentified.
XX
PM US2003092046-A1.
XX
PD 15-MAY-2003.
XX
PF 20-SEP-2002; 2002US-00247893.
XX
PR 07-JUL-1999; 99US-00349040.
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                invention may be useful for the detecting cytokeratin 18-20 expression an indicator for the diagnosis of tumour metastasis, particularly prostate cancer and lymphoma. The amplification using the primers is highly efficient and allows very sensitive detection of tumour metastasis. The current sequence is that of the human CK18-derived DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21-MAY-2002; 2002JP-00145689.
17-JUN-2002; 2002JP-00175271.
09-JUL-2002; 2002JP-00199759.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          human; cytokeratin; CK; LAMP; loop mediated isothermal amplification; tumour metastasis; prostate cancer; lymphoma; human; CK18; ss; primer probe; PCR; Flc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to novel nucleic acid amplification primers for the detection of human cytokeratin (CK) 18, 19 or 20 expression by the LAMP (loop mediated isothermal amplification) method. The primers of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LAMP nucleic acid amplification primers for detection of expression as indicator in diagnosis of tumour metastasis
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                                     20-SEP-2002; 2002US-00247893.
                                                                                                                                                                                    Duplex DNA strand 2 used for triplex formation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim
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                                                                                                                                                        diagnosis; therapeutic; investigation; ds
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                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.4; DB 1
Pred. No. 8.8e+02
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to novel oligonucleotides comprising several nucleotide units which are specifically hybridisable with a selected sequence of RNA or DNA wherein at least one of the nucleotide moieties of the oligomer is modified to include a guanidinium group. These oligonucleotides are useful for diagnostic, therapeutic and investigative purposes. The present sequence is a duplex DNA strand 1 used for triplex formation in the exemplification of the invention.
    The invention relates to a compounds. The guanidinium
                                                                                                                                                                                                                                                                                                                                                               07-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                              07-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ss; guanidinium
2-0-guanidinium
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(COOK/)
(PRAK/)
                                                                            Example 36; SEQ ID NO 10;
                                                                                                                                                   oligomers used for diagnostic,
                                                                                                                                                                                                                     WPI; 2004-118052/12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2004-031184/03
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PRAKASH T P.
MOHAN V.
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                                                                                                                                                                                                                                                                    Prakash
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    guanidinium functionalised nucleotide functionalised nucleotide compounds as
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Pred. No. 8.8e+02;
                                                                                                                                                nucleotide compounds useful for preparing therapeutic and investigative
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affinity; triplex formation
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This invention describes novel oligonucleotides derived from microsatellite markers and used for the amplification of the rose genome. The invention also describes a test kit for genetic analysis of cultured cor wild forms of the genus Rosa sp. that contains at least one of the new oligonucleotide primers and preparing microsatellite markers of Rosa sp. to PCR amplification of hypervariable genomic regions, using at least one primer pair, to produce polymorphic fragments which are separated and contents. The amplified markers are separated by electrophoresis, complified. The amplified markers are separated by electrophoresis, complified. The amplified markers are separated by electrophoresis, complicated the detected by staining (ethidium bromide or silver), complicated edetected by staining (ethidium bromide or silver), complicated with dyes or fluorophores or by mass spectrometry. A genomic library of 0.5-1.5 kb fragments from the rose variety conformation and the cells tested against a high-density array of synthetic coll and the cells tested against a high-density array of synthetic microsatellites. The eligonucleotides are used for genetic coll and the cells tested against a high-density array of synthetic conformations varieties. The oligonucleotides are used for genetic coll analysis of cultivated and wild types of roses, particularly for genetic constructed and wild types of roses, particularly for genetic constructed and wild types of roses, particularly for genetic constructed and wild types of roses, particularly for genetic constructed and wild types of roses, particularly for genetic constructed and wild types of roses, particularly for genetic constructed and wild types of roses, particularly for genetic constructed and wild types of roses, particularly for genetic constructed and wild types of roses, particularly for genetic constructed.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        microsatellite marker; rose genome; PCR; hypervariable region; genetic mapping; relatedness analysis; hybrid identification; breeding; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Page 9; 52pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New oligonucleotides from rose microsatellite markers, useful for genomic analysis, including identification of varieties and hybrids.
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No. 8.8e+02;
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The present invention is based in part on the discovery of novel secreted and membrane-bound polypeptides and their encoding polynucleotides. The nucleic acids and polypeptides are collectively referred as NOVX. The invention is useful for treating, preventing and diagnosing diseases such as metabolic disorders, diabetes, obesity, infectious diseases such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Parkinson's disease; epilepsy; immune disorder; osteoarthritis; haematopoietic disorder; inflammatory skin disorder; asthma; dyslipidemia; neurogenesis; cell differentiation; cell proliferation; haematopoiesis; wound healing; angiogenesis; chromosome mapping; tissue typing; preventive medicine; pharmacogenomic; gene therapy; anorectic; cardiant; virucide; antibacterial; fungicide; protozoacide; nootropic; neuroprotective; dermatological; human; PCR; primer; ss.
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                                                            The present invention relates to antisense compounds, compositions and methods used for modulating the expression of haem oxygenase (HO) 1. The methods and compositions of the present invention are useful for the diagnosis, prevention and/or treatment of diseases or conditions associated with aberrant expression or activity of haem oxygenase 1 such as hyperbilirubinaemia, neonatal jaundice and neurodegenerative diseases like Alzheimer's and Parkinson's disease. The invention is also useful in antisense-therapy. The present sequence is human haem oxygenase 1 DNA amplifying PCR primer used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of oxygenase 1, such as hyperbilirubinemia, neonatal jaundice and neurodegenerative diseases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Haem oxygenase 1; HO; hyperbilirubinaemia; neonatal jaundice; neurodegenerative disease; Alzheimer's disease; Parkinson's disease antisense-therapy; nootropic; neuroprotective; human; PCR; primer;
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                                                                                                                                                                     The present sequence is the antisense strand of a short interfering RNA (sirNA) targeted to nucleotides 1225-1243 of human insulin-like growth factor binding protein-5 (IGFBP-5) cDNA. The sense strand is also provided ADL70482. The sirNA can be used to interfere with the expression of IGFBP-5. Inhibition of IGFBP-5 expression can delay the progression of hormone-regulated (prostatic or breast) tumour cells to hormone-regulated cancers, provide a method for the treatment of hormone-regulated cancers, and inhibit or delay the growth and metastatic progression of prostate, breast and other IGF-1 sensitive tumours in bone. siRNAs of the invention can be used alone or in combination with other chemotherapy or apoptosis inducing treatments for the treatment of prostate cancer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jansen B,
Gonos ES;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New RNA molecule less than 49 bases and having a sequence effective to mediate degradation or block translation of mRNA that is the transcriptional product of a target gene, useful for treating Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         21-AUG-2002; 2002US-0405193P.
03-SEP-2002; 2002US-0408152P.
20-MAY-2003; 2003US-0472387P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RNA interference; RNA; short interfering RNA; siRNA; human; insulin-like growth factor binding protein-5; cytostatic; neuroprotective; nootropic; gene silencing; IGFBP-5; DNA-RNA hybrid; ss.
                                                                                         sarcomas such as osteosarcoma, renal cell carcinoma, breast cancer, bladder cancer, lung cancer, colon cancer, ovarian cancer, anaplast large cell lymphoma and melanoma, and also for the treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    disease or cancer.
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                                                                Alzheimer's
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 28; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGTTTGAGACAGCTGCC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gleave ME,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Signaevsky M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
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Pred. No. 8.8e+02;
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Best Local
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16-MAR-2000; 2000US-0189810P.
22-MAR-2000; 2000US-0191158P.
30-MAR-2000; 2000US-0193086P.
03-MAY-2000; 2000US-0201388P.
31-MAY-2000; 2000US-00584411.
The invention relates to human NOVX polypeptides and polynucleotides. NOVX sequences are useful in the treatment of cancer, hyperproliferative diseases, cirrhosis, keloid, psoriasis, tissue hypertrophy, osteoarthritis, atherosclerotic plaque formation, haemorrhage, ischaemic heart or renal disease, thrombosis, diabetes mellitus, hypertension, hypothyroidism, asthma, burns, incisions, ulcers, periodontal disease, Alzhemier's disease, Parkinson's disease, Huntington's disease, Alzhemier's disease, Parkinson's disease, Huntington's disease, amyotrophic lateral sclerosis, Shy-Drager syndrome, immune deficiencies and disorders such as severe combined immunodeficiency (SCID), bacterial infection, viral infection e.g. malaria, fungal infection e.g. candidiasis, autoimmune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               atherosclerotic plaque formation; haemorrhage; ischaemic heart disease; renal disease; thrombosis; diabetes mellitus; hypertension; hypothyroidism; severe combined immunodeficiency; SCID; infection; malaria; candidiasis; autoimmune disorder; connective tisque disease; multiple sclerosis; systemic lupus erythematosus; rheumatoid arthritis; autoimmune pulmonary inflammation; Guillain-Barre syndrome; autoimmune inflammatory eye disease; aravis; graft- versus-host disease; autoimmune inflammatory eye disease; arsthma; burn; incision; ulcer; periodontal disease; Alzhemier's disease; Parkinson's disease; Huntington's disease; amyotrophic lateral sclerosis; Shy-Drager syndrome; haematopoiesis; wound healing; tissue repair; antitumor; antitifalammatory. Drs. vience:
                                                                                                                                                                                                                                                                                                                                                                                                                                sclerosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel substantially purified NOVX polypeptide for treating severe combined immunodeficiency, candidiasis, cancer, asthma, multiple
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2004-068928/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Shimkets RA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-JUL-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; NOVX protein; cancer; hyperproliferative disease; keloid; psoriasis; tissue hypertrophy; osteoarthritis;
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No. 8.8e+02;
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RESULT 798
AAQ89896/c
ID AAQ898
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                                                      Query Match
Best Local S
Matches 16
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Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        lupus erythematosus, rheumatoid arthritis, autoimmune pulmonary inflammation, Guillain-Barre syndrome, autoimmune pulmonary dependent diabetes mellitus, myasthenia gravis, graft-versus-host disease and autoimmune inflammatory eye disease. The invention is useful in regeneration and tissue growth of bone, cartilage, tendon, ligament, haematopoiesis regulation, wound healing and tissue repair. Sequences of the invention also exhibit antitumor and antinflammatory activities. The present sequence is human NOVX DNA specific PCR primer used in the exemplification of the invention.
                                                                                                                                         AAQ89895 and AAQ89896 are a pair of PCR primers used in the detection of the cystic fibrosis mutant region G 542X, using a novel method of detection. The G 542X mutation features a G to T mutation at nucleotide position 1756 in exon 11 of the CPTR gene. The method is useful in detecting genetic mutations or abnormalities such as point mutations and detections and is therefore useful in the diagnosis of genetic and particularly inheritable genetic diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cystic fibrosis transmembrane conductance regulator; CFTR; mutational analysis; deletion; genetic abnormality; detection;
                                                                                                                Sequence
                                                                                                                                                                                                                                                                                          mutation and
                                                                                                                                                                                                                                                                                                          Detection of
                                                                                                                                                                                                                                                                                                                                      WPI; 1995-178882/23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9511990-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cystic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-JAN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ89896 standard; DNA; 22 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21 BP; 8 A; 4 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                             Example 1; Page 19; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                   Martinazzo
                                                                                                                                                                                                                                                                                                                                                                                                                           29-OCT-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-MAY-1995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disorders such as connective tissue disease, multiple sclerosis, systemic
                                                                                                                                                                                                                                                                                                                                                                                              (RAGG-) RAGGIO-ITALGENE SPA.
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                           4876 TITCTTTCTCTGCAAAC 4892
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21
                                                        16;
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                                                                      Similarity
 TGTCI
                                                                                                               22 BP; 10 A; 4 C; 5 G; 3 T; 0 U; 0 Other;
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                                                        Conservative
                                                                                                                                                                                                                                                                                        genetic mutation(s) - using one probe to one side second and third probes spanning the mutation.
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                                                                                                                                                                                                                                                                                                                                                                   Bichi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        94WO-EP003538
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chromosome 7 mutant
                                                                    0.3%;
                                                                                                                                                                                                                                                                                                                                                                 'n
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                                                                                                                                                                                                                                                                                                                                                                 Marcolini S,
                                                      Sccre 15.4; D
Pred. No. 8.8e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       No. 8.8e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     region,
                                                                                                                                                                                                                                                                                                                                                                 Turchetti E,
                                                                    8.8e+02
                                                                                     DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     542X PCR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                  Length
                                                        Indels
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                                                                                                                                                                                                                                                                                                                                                                   Pelliccia
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RESULT 799
RAT78996/c
ID AAT789
XX AAT789
XX Human
XX Human
XX Human
XX Homo E
XX CA2178
PM CA2178
PM CA2178
PM O1-JUN
XX (UYBR-
XX WPI;
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XX WPI;
I Hayder
XX Mouse
PT As a |
XX Mouse
PT As a |
XX This
CC This
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CC Gene,
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ARASULT 800
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XX ARAS66
XX ARAS66
XX COBPIE
XX COUPLE
XW CESTIL
XW Waste
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Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This oligonucleotide comprises the 5' acceptor site of intron 1 of the human Huntington's disease (HD) gene. The splice site sequences for the first 5 exons of the mouse HD gene (see AAT78974) and the human HD gene were compared (see AAT78985-T79002). Targeted disruption of the murine HD gene, e.g. at exon 5, can be used to examine the function of the HD gene and its role in development. Transgenic mice can be used as models of HD
                                                                                                                                                           Coupled polymerase chain reaction; PCR; ligase detection rearestriction endonuclease digestion; RED; infectious disease; waste water purification; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse Huntington's Disease gene - useful for generating transgenic mice as a model of Huntington's Disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human Huntington's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT78996;
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                                                       WO200056929-A2
                                                                                                            Unidentified
                                                                                                                                                                                                                                                                     Primer used in mismatch extension of DNA in LDR reactions.
                                                                                                                                                                                                                                                                                                                            08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                              AAA96617;
                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA96617 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 60; 69pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hayden M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-JUN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYBR-) UNIV BRITISH COLUMBIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Lin B, Nasir J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 2 A; 3 C; 1 G; 16 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first
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                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             disease
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gene; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ₽₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.4;
Pred. No. 8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22
                                                                                                                                                                                                                     reaction; LDR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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AALSOLT 801
AALSO570/c
ID AALSO5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The specification describes the use of coupled polymerase chain reaction (PCR), restriction endonuclease digestion (RED) and ligase detection reaction (LDR) to identify one or more low abundance sequences differing by one or more single base changes, inserts or deletions, from a high abundance sequence, in several target nucleotide sequences. The method is used to identify one or more low abundance sequences. The method is used to identify one or more low abundance sequences. The method is also useful for detecting a wide variety of infectious diseases caused by bacterial, viral, parasite and fungal infectious agent. Cancers can also be detected by this method. The method is also used for detection, identification and monitoring of pathogenic and indigenous municipal waster water purification system and water reservoirs or in polluted areas undergoing bioremediation and to detect plasmids containing genes that can metabolise xenobiotics, to monitor specific target microorganisms in population dynamic studies, or either to detect, identify, or monitor specific target microorganisms modified microorganisms in the environment and in industrial plants. The present sequence represents a primer used in mismatch extension, in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local S
Matches 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Coupled polymerase chain reaction-restriction endonuclease digestion-
ligase detection reaction to identify low abundance sequences differing
by single-base changes, insertion or deletion from high abundance
sequence in target sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (CORR )
 16-MAR-2001; 2001GB-00006635.
02-AUG-2001; 2001GB-00018879.
                                                                                                                                                                                                Molecular array;
gene expression;
                                                                                                                                                                                                                                                    Molecular array
                                                                                                                                                                                                                                                                                                                                                              AAL50570 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Barany F,
                                                      18-MAR-2002; 2002WO-GB001245
                                                                                                                            WO200274988-A2
                                                                                                                                                                Unidentified
                                                                                                                                                                                                                                                                                         12-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4737 GGAGACCCATCCTCACC 4753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                      GGAGGCCCATCCTCACC 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22 BP; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Fig 6C; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Day JP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
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                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                    production method-related PCR primer.
                                                                                                                                                                                                   SNP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%; Score 15.4; DB 1;
94.1%; Pred. No. 8.8e+02;
                                                                                                                                                                                              target molecule identification; genetic analysis; detection; haplotyping; sequencing; PCR; primer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                9 C; 4 G; 4 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention comprises a method for producing a molecular array, the method involves immobilising molecules to a solid phase at a density which allows individual immobilised molecules to be individually resolved. The molecular array produced by the method of the invention useful for identifying one or more target molecules in a sample. The molecular array is also useful in genetic analysis, gene expression studies, identifying molecules which interact with a target molecule, detection/typing of single nucleotide polymorphisms, haplotyping and sequencing. The present DNA sequence represents a PCR primer that was used in an example of the invention
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Pred. No. 8.8e+02
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The present sequence is that of pyrene-anchored locked nucleic acid (INA) CC oligo(dT) primer ON14, which was used in first-strand cDNA synthesis from CC eukaryotic mRNA. It includes compound '17d' at its 3' end, which is based CC on an INA-type 2'-0,4'-C-methylene-beta-D-ribofuranosyl molety. It is CC one of a set of such primers (see also ACC48482-85) that were used in an CC example from the invention to demonstrate improved reverse transcription CC of mRNA using pyrene-LNA anchored oligo(T) primers. The following results CC were observed: efficient priming on mRNAs with short poly(A) tails; CC efficient anchoring of the oligo(T) primer by pyrene-LNA and LNA-C/G/T CC units resulting in an improved T20-VN anchor primer and thus avoiding CC reverse transcription of long poly(A) tracts; and improved reverse conscription of eukaryotic poly(A) tracts; and improved reverse concreased specificity. The invention relates to modified LNA units CC that comprise unique base groups. Desirable nucleobase and nucleosidic CC base substitutions can mediate universal hybridisation when incorporated into nucleic acid strands. The novel LNA compounds can be used e.g. as CC and in diagnostics.
                                                                                                                                                                                                                                                                                                       Novel nucleic acid comprising a locked nucleic acid unit having a modified base that comprises an optionally substituted carbocyclic moiety, or modified nucleobase or nucleosidic base other than
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22-SEP-2001; 2001US-0323967P.
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coligo(GT) primer ONIS, which was used in first-strand CDNA synthesis from CCC oligo(GT) primer ONIS, which was used in first-strand CDNA synthesis from CCC enkaryotic mRNA. It includes compound '17d' at its 3' end, which is based CC one of a set of such primers (see also ACC48482-84) that were used in an CCC example from the invention to demonstrate improved reverse transcription of mRNA using pyrene-LNA anchored oligo(T) primers. The following results were observed: efficient priming on mRNAs with short poly(A) tails; CC were observed: efficient priming on mRNAs with short poly(A) tails; CC units resulting in an improved T20-VN anchor primer and thus avoiding reverse transcription of long poly(A) tracts; and improved reverse transcription of eukaryotic poly(A) tracts; and improved reverse conscription of eukaryotic poly(A) tracts; and improved reverse that comprise unique base groups. Desirable nucleobase and nucleosidic that comprise unique base groups. Desirable nucleobase and nucleosidic consensation when incorporated into nucleic acid strands. The novel LNA compounds can be used e.g. as CCC PCR primers, in sequencing, the synthesis of antisense oligonucleotides.
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Matches 17; Conserv
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22-SEP-2001; 2001US-0323967P.
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RESULT 804
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IDC484483/c
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Best Local (
                                                                     04-SEP-2001;
22-SEP-2001;
                                                                                                                        04-SEP-2002; 2002WO-IB003911.
                                                                                                                                                                                              WO2003020739-A2.
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 Wengel J,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Locked nucleic acid anchored oligo(I) primer ON13.
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                                  (EXIQ-) EXIQON AS
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                                                                     2001US-0317034P.
2001US-0323967P.
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/mod_base= OTHER
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22
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/mod_base= OTHER
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mod_base= OTHER
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Pred. No. 8
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    Indels

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The present sequence is that of pyrene-anchored locked nucleic acid (LNA) CC oligo(dT) primer (NN3, which was used in first-strand cDNA synthesis from CC eukaryotic mRNA. It includes compound '17d' at its 3' end, which is based CC on an LNA-type 2'-O,4'-C-methylene-beta-p-riboturanosyl molety. It is CC one of a set of such primers (see also ACC48482-85) that were used in an CC example from the invention to demonstrate improved reverse transcription CC were observed: efficient priming on mRNAs with short poly(A) tails; CC efficient anchoring of the oligo(T) primers The following results CC units resulting in an improved T2O-VN anohor primer and thus avoiding CC reverse transcription of long poly(A) tracts; and improved reverse transcription of eukaryotic poly(A) tracts; and improved reverse CC transcription of eukaryotic poly(A) +RNA directly from total RNA extracts CC text comprise unique base groups. Desirable nucleobase and nucleosidic consess substitutions can mediate universal hybridisation when incorporated conto nucleic acid strands. The novel LNA compounds can be used e.g. as CC primers, in sequencing, the synthesis of antisense oligonucleotides, and in diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel nucleic acid comprising a locked nucleic acid unit having modified base that comprises an optionally substituted carbocycl moiety, or modified nucleobase or nucleosidic base other than oxazole/imidazole.
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밁 S Query Match
Best Local Similarity
Matches 17; Conserv 5392 TAAAAAATACAAAAAAGAAA 5412 ;||||||| | ||||| ||| 21 BAAAAAAAAAAAAAAAAA 1 0.3%; llarity 81.0%; Conservative ۲. Score 15.4; DB 1; Pred. No. 8.8e+02; 1; Mismatches 3 Length Indels 22; 0 Gaps 0

Sequence 22 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 2 Other;

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RESULT 805
ABZ80983
ID ABZ809
XX
AC ABZ809
XX
DT 15-OCT
XX
DW MOUSE
XX
DW MOUSE
XW BEST:
KW ANTILL
KW ANTILL
KW HOER
XW BLEE
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PN WO2(
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PR 09-
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PR 09-
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                                                                                                                                                                                                                                                                                    Bestrophin; vitelliform macular dystrophin 2; VMD2; ss; PCR; vaccine; antiobesity; antidiabetic; immunomodulator; hypotensive; cardiant; antilipemic; osteopathic; antiinflammatory; cytostatic; obesity; primer energy homeostasis; metabolism; triglyceride; body-weight regulation; eating disorder; cachexia; diabetes mellitus; hypertension; gallstones;
                                                                                                                                                                                                                                                         BIEED apnea.
                                                                                                                                                                                                                                                                                                                                                                                    Mouse vitelliform macular dystrophin 2-like protein 3 gene primer #2.
                                                                                                                                                                                                                                                                                                                                                                                                                     15-OCT-2003
                                                                                                                                                                                                                                                                          coronary heart disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                        hypercholesterolemia; osteoarthritis;
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primer;

WPI; 2003-393411/37. Steuernagel A, 09-OCT-2001; 2001EP-00124059 09-OCT-2002; 2002WO-EP011321.

DEVELOGEN ENTWICKLUNGSBIOLOGISCHE FORSCH

Broenner G,

Fritsch R,

Eulenberg

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Ciossek

Mus musculus.

17-APR-2003 WO2003030922-A2

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RESULT 806
RESULT 806
RAD51324/c
ID AAD513
XX AAD513
XX AAD513
XX IO 16-APR
XX Anchor
XX Lamini
KW Gastri
KW Gastri
KW Gastri
KW WO2002
XX Uniden
XX 
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Best Local S
Matches 16
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                                                                                                                                                                                          Assessing condition e.g. athletic ability, stage of disease, presence of drugs, response to exercise, response to vaccines, therapies, nutritional states, of performance animal involves analyzing nucleic acid expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             gastritis; gastric ulcer; r
musculoskeletal damage; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 22 BP; 5 A; 6 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New pharmaceutical compositions comprising a Bestrophin gene, polypeptide or nucleic acid, for treating, alleviating and/or preventing metabolic diseases, e.g. obesity, cachexia, diabetes mellitus, hypertension, or
                                                                                                                                                                                                                                                                                                                                                                                       Brandon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             04-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           03-MAY-2002; 2002WO-AU000553
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Laminitis; viral disease; vaccine; bacterial disease; primer; epistaxis; gastritis; gastric ulcer; respiratory ailment; fracture; joint disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Anchored oligo dT primer used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAD51324 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 4; Page 49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-APR-2003
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2001US-00896941.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  PARTNERS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP
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Pred. No. 8.
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performance of expressed

Disclosure; Page 46;

87pp; English

The invention relates to a method for assessing a condition of a performance animal. The method involves determining in sample about of expressed target nucleic acid; transmitting digital sample signals.

sample signal

abundance signal to

Sequence 22 illustrate the

BP;

method 0 A; 0 C;

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Other;

homology search on a database. The methods are useful for presentation of sequences. It is also useful to prepare primer sequences to obtain a cDNA clone of a total coding region from a partial sequence of a gene having an unidentified sequence. The present sequence is a primer specific for human chromosome 10 RP-11-336AlO clone DNA. This sequence is used to

The invention relates to methods and system for sequence presentation. The method involves extracting a partial sequence corresponding to a partial sequence of a target gene having an unidentified sequence, by

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AAD64451/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Presenting partial sequences by predicting and extracting exon sequences from a database, is useful to prepare primers to obtain a cDNA clone of total coding region from a partial sequence of an unidentified gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human RP-11-336A10 clone specific primer
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Pred. No. 8.
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27-DEC-2001;

27-DEC-2001;

31-DEC-2001;

11-MAY-2002;

15-MAY-2002;

28-MAY-2002;

29-MAY-2002;

29-MAY-2002;

29-MAY-2002;
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                        New NOVX polypeptides and nucleic acids, useful for preventing or treating NOVX-associated disorders, e.g. cancer, cardiomyopathy, atherosclerosis or diabetes, and in chromosome mapping, tissue typing
                                                                                                                                                                              Alsobrook JP, Anderson DW, Boldog FL, Burgess CE, Chillakuru Eddinger SR, Gerlach VL, Gorman L, Gould-Rothberg BE, Guo X; Jeffers ME, Ji W, Li L, Malyankar UM, Miller CE, Murghey R; Patturajan M, Peyman JA, Rastelli L, Rieger DK, Shenoy SG; Smithson G, Starling G, Taupier RJ, Voss EZ, Zhong H, Zhong
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2001US-0341346P.
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2001US-0344297P.
2002US-0344903P.
2002US-0380981P.
2002US-0381495P.
2002US-0383534P.
2002US-0383534P.
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2002US-0383539P.
2002US-0406353P.
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; Pred. No. 8.8e-
1; Mismatches
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CC antiarteriosclerotic, hypotensive, cytostatic, anorectic, antirheumatic, communosuppressive, antidiabetic, nephrotropic, dermatological, communosuppressive, anti-HIV, antiinflammatory, neuroprotective, cc nootropic, antipsoriatic, antiparkinsonian, antiasthmatic, neuroleptic, cc notropic, antipsoriatic, antiparkinsonian, antiasthmatic, neuroleptic, cc antidepressant, antiastlergic or gynaecological activities. The DNA cc sequences of the invention may be useful for gene therapy whilst the cc protein sequences may allow the development of a vaccine. The protein is cuseful in the manufacture of a medicament for treating a syndrome cc useful in the manufacture of a medicament for treating a syndrome cc diagnosing, treating or preventing NOVA-associated disorders, for example cc ardiomyopathy, atherosclerosis, hypertension, cancer, obsetly, cr heumatoid arthritis, diabetes, glomerulonephritis, psoriasis, skin cc disorders, AIDS, inflammation, multiple sclerosis, Alzheimer's disease, extina, schizophrenia, depression, allergies or fertility disorders. The nucleic acids may further be used as the hybridisation probes, in chromosome mapping, tissue typing, preventive medicine, and pharmacogenomics. The present sequence is that of PCR cc medicine, and pharmacogenomics. The present sequence is that of pcR cc medicine, and pharmacogenomics seed for the amplification of a human cc sequence during the analysis of gene expression in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                encode them, having properties related to stimulation of blochemical or physiological responses in a cell, a tissue, an organ or an organism. Compounds which modulate the proteins of the invention may have cardiant,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention
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Sequence 22 BP; 3 A; 6 C; 3 G; 10 7 0 U; 0 Other;

뫉 8 Best Matches Local 2942 Similarity CCAGAAACCTGAAGGAG 2958 Conservative 0.3%; 4 0; Score 15.4; Pred. No. 8 Mismatches .8e+02; DB 1; Length 22; Indels 0, Gaps 0

ADF88323
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AC ADF
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OS RESULT 809 JP2003235571-A. Synthetic. human; single nucleotide polymorphism; microarray; side effect; ss; primer; PCR. Single nucleotide polymorphism detection primer, SEQ ID No 1906 26-FEB-2004 ADF88323 ADF88323 standard; Homo sapiens. (first DNA, entry) 22 ВP

26-AUG-2003

12-FEB-2002; 2002JP-00034717.

12-FEB-2002; 2002JP-00034717

(KAGA-) KAGAKU GIJUTSU SHINKO JIGYODAN

WPI; 2003-820454/77.

Novel polynucleotide in human gene. useful for detecting single nucleotide polymorphisms

Claim 2; SEQ ID NO 1906; 704pp; Japanese

The invention relates to a novel polynucleotide isolated and purified

99999999999999998**%**

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RESULT 810
ABX74887/c
ID ABX748
XX ABX748
XX ALX748
XX A
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Best Local Sim:
Matches 16;
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                                The present invention relates to a microarray comprising a matrix of at least one cDNA probe from a set of probes immobilised to a solid surface in a predetermined order, where a row of pixels corresponds to replicates of one distinct probe from the set. The probes are complementary to nucleic acid sequences that are expressed differentially in aggressive as compared to non-aggressive types of clear cell renal carcinoma (CC-RCC) and that hybridise to the probes under high stringency conditions. The microarray is useful for the prognosis of patients with CC-RCC, wherein aggressive and non-aggressive CC-RCC tumour types are characterised by aggressive and non-aggressive CC-RCC tumour types are characterised by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    differential expression profile; aggressive CC-RCC tumour type; non-aggressive CC-RCC tumour type; clear cell renal carcinoma; gene expression profiling; tumour tissue; oligo-dT; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New microarray, comprising a matrix of cDNA probe from immobilized to a solid surface in predetermined order, prognosis of patients with clear cell renal carcinoma.
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            aggressive a
differential
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 30; 179pp; English.
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Pred. No. 8.8e+02;
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useful in the
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                                         The invention relates to a method of selecting genes expressed in a CC cancer cell, which involves characterising cancer based on the CC cancer cell, which involves characterising cancer based on the CC functioning of gene selection by comparing the expression of the selection CC gene from the cancer cell with the expression of an identical selection CC genes from a noncancerous cell or different type of cancer cell. The CC method is useful for targeting the therapy of cancer by using a selection CC method is useful for targeting the therapy of cancer by using a selection of genes or their products expressed in a cancer cell, the gene selection CC comparing the expression of the selected gene or their products from the cancer cell with the expression of an identical selection of genes or their products noncancerous. The method is also useful for diagnosing, their products monitoring and classifying a disease condition e.g. cancer ccl such as neuroblastoma, rhabbomyosarcoma, Burkitt's or Ewing family of tumours. The present sequence represents an anchored oligonucleotide
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Best Local S
Matches 17
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Ewing
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                                                                                                                                                                                                                                                                                                                         Selecting genes expressed in cancer cell, by characterizing cancer based on functioning of gene selection by comparing expression of selected gene from cancer cell with expression of selected genes from noncancerous
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                                                                                                                                                                                                                                                                                  Example
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17; Conserv
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RINGNER M.
PETERSON C.
MELTZER P.
                                                                                                                                                                                                                                                                                  2; Page 18; 53pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                  Ringner M,
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Pred. No. 8.8e+02;
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06-MAY-2004

(first

entry)

Human fibrosis/scarring predisposition-related PCR primer SeqID25.

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RESULT 812
ADX69815/c
ID ADX698
XX ADX698
XX ADX698
XX O6-MAY
XX PCR; p
KW Chondr
KW D-gluc
XX DF2004
XX JP2004
XX J
RESULT 813
ADJ45826
ID ADJ458
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention relates to a novel glucosyl-transfer agent. Specifically, it refers to the synthesis of chondroltin or chondroltin-sulphate proteoglycans using this glucosyl-transfer agent to transfer the N-acetyl galactosamine residue from a donor substrate to D-glucurone acid existing in the non-reducing terminal of an N-acetyl-D-galactosamine receptor substrate containing a formulaic sugar chain. The present invention describes a detection method for the canceration of tissue associated with the glucosyl-transfer agent as CSGalNac-T. This oligonucleotide sequence is a PCR primer given in an exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Novel glucosyl-transferase transferring N-acetyl-D-galactosamine with respect to linkage tetrasaccharide existing in binding site chain, useful for synthesizing chondroitin/chondroitin-sulfate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PCR; primer; human; glucosyl-transfer agent; chondroitin; chondroitin-sulphate proteoglycan; N-acetyl galactosamine; D-glucurone acid; canceration; CSGalNAc-T; ss.
                                                                                    ADJ45826 standard; DNA; 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 22 BP; 2 A; 9 C; 4 G; 7 T; 0 U; 0 Other;
                              ADJ45826
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 1;
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Pred. No. 8.8e+02
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in vitro diagnosis; inappropriate fibrosis; scarring; 16sRNA region; ND2 gene; NADH-quinone oxidoreductase; complex I; cytochrome b; complex III region; COI gene; Cytochrome coxidase; complex IV region; mitochondrial genome; antiinflammatory; vulnerary; dermatological; nephrotic; gene therapy; Dupuytren's disease; keloid; hypertropic scar; scleroderma; systemic sclerosis; crest syndrome; tuberous sclerosis; skin patch; familial cutaneous collagenoma; skin metabolic disorders; eosinophilic fascitis; discoid lupus erythematosus; dermatomyositis; mixed connective tissue disease; drug-induced skin fibrosis; oral submucous fibrosis; pulmonary; cardiac fibrosis; liver fibrosis; cirrhosis; renal fibrosis; drug induced fibrosis; liver fibrosis; central nervous system fibrosis; peripheral nervous system fibrosis; gynaecological fibrosis; genitourinary tract fibrosis; gynaecological fibrosis; glomerulonephritis; cystic fibrosis; scleroderma; myocardial fibrosis; myocardial infarction; stroke; neurodegenerative disorder; human; PCR; primer; ss.
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Homo sapiens.

02003093506-A2.

13-NOV-3003

23-APR-2003; 2003WO-GB001717

30-APR-2002; 2002GB-00009812

(RENO-) RENOVO LTD.

Ferguson MWJ, Ollier WER, Bayat A;

WPI; 2004-022664/02.

In vitro diagnosing a condition due to inappropriate fibrosis or scarring, e.g. Dupuytren's disease by detecting polymorphism or mutation in the 16sRNA region, ND2 gene, cytochrome b region or COI gene of the mitochondrial genome.

Example 1; SEQ ID NO 25; 81pp; English

This invention relates to a novel method of in vitro diagnosis or condetction of a predisposition to a condition characterised by composition for predisposition to a condition characterised by the 16sRuh (C region, ND2 gene of NDH-quinone oxidoreductase (complex I), cycochrome by (complex III) region or COI gene of Cytochrome c oxidase (complex IV) (C region of the mitochondrial genome to detect the presence of a genetic polymorphism or mutation linked to the development of the condition. The continuous proves useful for the development of the condition. The continuous proves disclosed may prove useful for gene therapy. The caddition, the sequences disclosed may prove useful for gene therapy. The continuous predisposition to a condition characterised by inappropriate c fibrosis or scarring. A modulator of mitochondrial genome gene products is useful in the manufacture of medicament for treating a condition of characterised by fibrosis or scarring. The condition is Dupuytren's case syndrome, tuberous sclerosis with skin, ecsinophilic fascitis, collagenome, and the skin fibrosis, oral submucous fibrosis, fibrosis induced skin fibrosis, oral submucous fibrosis, fibrosis induced skin fibrosis, oral submucous fibrosis, fibrosis collowing a stroke or scarcinal fibrosis, and central nervous system fibrosis, scentral nervous system fibrosis, fibrosis of collowing a stroke or neurodegenerative dissorders. The process of collowing a stroke or neurodegenerative dissorders. The process of collowing a stroke or neurodegenerative dissorders in the collowing a stroke or neurodegenerative dissorders.

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RESULT 814
ADL97794/c
ID 97794/c
ID 01977
XX ADL977
XX BD1977
XX BS; pr
Oligon
XX BS; pr
OS Uniden
XX W02004
XX W1-APR
PF 19-SEF
PR 19-SEF
PR 19-SEF
PR 19-SEF
PR 19-SEF
XX UYCH-
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PI Produc
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                                                                                                                                                                                                                              The invention relates to a method of producing (M1) a molecular array, involves: immobilizing to a solid phase a several molecules at a density which allows individual immobilized molecules to be individually resolved, where each molecule in the array is spatially addressable and the identity of each molecule is known or determined prior to immobilization; and optionally providing a molecular array comprising a several molecules immobilized to a solid phase at a density such that individual immobilized molecules are not capable of being individual resolved, and reducing the density of functional immobilized molecules in the array such that remaining individual functional immobilized molecules in the array such that remaining individual functional immobilized molecules are capable of being individually resolved, where each individual functional molecule in the resulting array is spatially addressable and the identity of each molecule is known or determined prior to the density reduction step. The array efficiently resolve complex samples, separate correct signals from erroneous signals, eliminates need for sample amplification, detects transient interactions or temporal characteristic amplification to the correct stransient interactions or temporal characteristic and incomplete the corrections.
Best Loc
Matches
                                                         Query Match
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Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Producing molecular array by immobilizing to solid phase several known molecules at low density for allowing individual immobilized molecules be individually resolved and spatially addressable.
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                                                                                                                        Sequence 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 152; 219pp; English.
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26-SEP-2002; 2002GB-00022412.
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                               Local
                                                                                                                                                                                 single molecule processes. This sequence represents anoligonucleotide and in the method of the invention.
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                                                                                                                     BP;
Conservative
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                                                                                                                        0
                                                                                                                     A; 0 C; 0 G; 20 T; 0 U; 2 Other;
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                            81.0%;
                                                            0.3%;
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   7,
Pred. No. 8.8e+02;
1; Mismatches 3
                                                         Score 15.4;
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Pred. No. 8.8e+02;
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                                                         DB 1; Length 22;
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   Gaps
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RESULT 815
AAQ11215
ID AAQ112
XX AAQ112
XX AAQ112
XX DII Oligon
XX Beta 1
XX JP0304
XX JP0304
XX JP0304
XX JP0314
XX
RESULT 816
AAQ10290
ID AAQ102
XX
AC AAQ102
XX
DT 05-APR
XX
Probe
XX
Probe
XX
Polyme
XX
Synthe
XX
PN EP4091
XX
PD 23-JAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligo-nucleotide for detecting Bacillus cereus in specimen - has sequence complementary to sequence coding for B cereus beta-lactamase gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This oligonucleotide is used as a primer for a polymerase chain reaction (PCR), in conjunction with other primers (see AAQ11212-14 and AAQ11216), to selectively amplify a target sequence prior to its detection by electophoresis or chromatography. It has a seq- uence complementary to a portion of the B.cereus beta-lactamase gene. It is useful in the detection of Bacillus cereus bacteria in samples, e.g. urine or food. Segmentation of the control of the cereus bacteria in samples, e.g. urine or food. Segmentation of the control of the cereus bacteria in samples, e.g. urine or food. Segmentation of the control of the cereus bacteria in samples, e.g. urine or food.
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                                                                                                                                                                                                                                                              Probe to beta-lactamase gene of Bacillus cereus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 1 A; 7 C; 3 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page 1; 6pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-MAR-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Beta lactamase; PCR; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ11215;
                                                                                        EP409159-A.
                                                                                                                                                Synthetic.
                                                                                                                                                                                                          Polymerase chain
                                                                                                                                                                                                                                                                                                                              05-APR-1991
                                                                                                                                                                                                                                                                                                                                                                                      AAQ10290;
                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ10290 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             also J0349697,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1991-106868/15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-JUL-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             JP03049696-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SHMA ) SHIMADZU CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5392 TAAAAAATACAAAAAAGAAA 5412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   _
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTCTGAAAGCCCATTCCCTG 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Trererareceerriceere 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BAAAAAAAAAAAAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -98 and -99
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    89JP-00185681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            89JP-00185681.
                                                                                                                                                                                                             reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                             PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                       food poisoning;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         e.g. urine or food. See
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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RESULT 817
AAQ25565/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-JUL-1989;
18-JUL-1989;
27-SEP-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ohashi T,
Aoyama Y,
                                                                                 New dye-coupled modified nucleosides, nucleotides and oligo:nucleotides useful for synthesis of antisense DNA and RNA strands in presence of template, also for in=vivo and in=vitro detection of genetic material.
                                                                                                                                                                                                                                                                                                                                               06-DEC-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dye-coupled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25-MAR-2003
02-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAQ25565 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 1 A; 7 C; 3 G; 9 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Probes may be used to identify food using the polymerase chain reaction
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Detection of bacterial species causing food poisoning chain reaction to amplify specific gene fragments.
                                      Example; Page 9; 17pp; German.
                                                                                                                                                                                                               Engels
                                                                                                                                                                                                                                                                                                      11-DEC-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                     EP490281-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA synthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ25565;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1991-023778/04.
                                                                                                                                                                      WPI; 1992-201578/25
                                                                                                                                                                                                                                                            (FARH ) HOECHST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-JUL-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SHMA ) SHIMADZU CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3; Page 53; 56pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 64 TTCTGAAAGCCCATTCCCTG 83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Treferarcecerrecere 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jikuya H,
Tada J;
                                                                                                                                                                                                               Herrlein M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3'-amino modified oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   89JP-00185685.
89JP-00185685.
89JP-00251400.
                                                                                                                                                                                                                                                                                                      90DB-04039488
                                                                                                                                                                                                                                                                                                                                               91EP-00120935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNA; antisense strands; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90EP-00113661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /*tag= a
/note= "3-amino modified"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Takano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
                                                                                                                                                                                                                   Konrad
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ç
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ģ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. 9.3e+02
                                                                                                                                                                                                                 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Shirasaki Y,
                                                                                                                                                                                                                 Mag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              poisoning agents
                                                                                                                                                                                                                 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Abe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ħ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yamagata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B.cereus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          polymerase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ã
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The sequence is an example of a

dye coupled 3'-amino modified oligo-

Query Match

Sequence

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A; 0 C; 0

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20 T; 0 U; 0 Other;

0.3%;

Score 15.2;

DB 1; Length 20;

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RESULT 818
AAQ33554/c
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문
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                                                         The sequence is that of a bovine microsatellite sequence obtd. by CC screening a library of bovine MooI DNA fragments of between 250 and 500 CC bp with an (AC)15 and a (TC)15 oligonucleotide probe. One out of 50 CC clones cross-hybridised. Assuming independent distribution of microsatellites and MooI sites, the frequency of (T6)n >9 microsatellites (T6)n >9 microsatellites and MooI sites, the frequency of (T6)n >9 microsatellites (T6)n >9 microsatellites (T6)n >9 microsatellites (T6)n >9 microsatellites (T7)n >9 mic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleotide, it can be used in the synthesis of DNA and RNA nucleosides, nucleotides and oligonucleotides and for the synthesis of opposite strands in the presence of a template strand and in fluorescence microscopic and macroscopic detection in vivo and in vitro of genetic material. It is labelled with a fluorescent dye. See also AAQ25566 and AAQ25567. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                       Table 7; Page 150; 517pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Polymorphic bovine DNA markers - used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Georges M, Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             06-AUG-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PCR; selection; pgenetic mapping;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Microsatellite sequence from clone AGLA247.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-MAR-2003
02-FEB-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAQ33554;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ33554 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1992-284684/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JAN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JAN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9213102-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (GENM-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5393 АДАДАДАТАСДАДАДАДАДА 5412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENMARK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     and selective breeding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.3%;
ilarity 85.0%;
Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  91US-00642342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primers;
; traits;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92WO-US000340.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPTIPRIM; breeding; amplification; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ₽P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             genetic identification,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cattle;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            parentage;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
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RESULT 819
AAQS8578
AC AAQ585
XX AAQ585
XX AAQ585
XX AAQ585
DT 25-MAI
DT 21-AU
XX Novel
XX Novel
XX Novel
XX W094
XX W994
XX I3-;
XX I1-
XX IN-
PR 11-
PR 11-
XX WP
XX UST
XX YFT WF
PT WF
PT WF
PT C1
XX YFT C1
XX CC CC
XX SQC
RESULT 820
AAQ94205/c
ID AAQ942
XX
AC AAQ942
XZ
DT 25-MAR
DT 24-AUG
DX
XX
XX
XX
XX
XX
XW Oligon
                                                                                                                                                                                                                                Matches
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                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                          The inventors claim a method of transporting a nucleic acid deriv. a membrane which comprises using a receptor that uses salt bridgin, aromatic stacking, H bonding and chelation to recognise the nucleic deriv. AAQ56305, AAQ58577-86 are nucleic acid derivs used in the examples. (Updated on 25-MAR-2003 to correct PN field.)
 Oligonucleotide ligand; steroid hormone; hapten; immobilisation;
                                                    25-MAR-2003
24-AUG-1995
                                                                                                                                                                                                                                                                                 Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                            Transport of nucleic acid derivs. across membranes - which use salt bridging, aromatic stacking, hydrogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                          Alpha-anomeric oligonucleotide ligand 1803 for oestradiol hapten
                                                                                                                                                                                                                                                                                                                                                                                     Example; Table 1, page
                                                                                                                                                                                                                                                                                                                                                                                                                 chelation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1994-082846/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-AUG-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-AUG-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-MAR-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9404194-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-MAR-2003
21-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAQ58578 standard; RNA; 20
                                                                                         AAQ94205;
                                                                                                                  AAQ94205 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ58578
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MASI ) MASSACHUSETTS INST TECHNOLOGY.
                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                5393 AAAAAAATACAAAAAAGAAA 5412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5393
                                                                                                                                                                                                                                 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 AAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAAAAAATACAAAAAAGAAA 5412
                                                                                                                                                                                 AAAAAAAAAAAAAAAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rebek J,
                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            synthetic RNA oligo which is a target nucleotide for a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                  (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93WO-US007603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        nucleic acid;
                                                                                                                                                                                                                                            0.3%;
85.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           De
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                85.0%;
                                                                                                                                                                                                                                                                                                                                                                                      38; 103pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mendoza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No. 9.3e+02;
0; Mismatches 3
                                                                                                                                                                                                                                                          Score 15.2;
                                                                                                                                                                                                                                                Pred.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        transport;
                                                                                                                                                                                                                                                No.
                                                                                                                                                                                                                                              .3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligo;
                                                                                                                                                                                                                                                          DB 1;
                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             using new receptors bonding and
                                                                                                                                                                                                                                                                                                                                                             acid deriv. accros
                                                                                                                                                                                                                                                          20;
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                                                                                                                                                                                                                                 Gaps
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                                                                                                                                                                                                                                                                                                                                     acid
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                                                                                                                                                                                                                                 0
RESULT 821
AAQ75585/c
ID AAQ755
XX AAQ755
XX AAQ755
XX CA-AUG
AC Revers
XX Revers
XX Revers
XX Analys
XW Analys
XW Analys
XW Synthe
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                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                Oligonuclectides (AAQ94201-Q94205) were synthesised for use as ligands. The ligands are covalently linked to a hapten (esp. a steroid hormone) to form a conjugate which is then immobilised on a solld support for interaction with antibodies against the hapten. Nucleic acid ligands are less likely to be recognised by the antibodies than are peptide ligands and nucleic acids are also less likely to undergo intramplecular organisation which interferes with accessibility of the hapten to the antibodies. For immunodiagnosis of oestradiol, the active hapten esstradiol-6-carboxymethoxime-N- hydroxypuctinimide ester was used.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Assay device for hapten or its specific antibodies - comprises support having competitive reagent immobilised via nucleic acid ligand to improprientation and accessibility.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cros
                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                         aggregate;
                                      Analysis;
                                                              Reverse transcription primer used in cDNA analysis technique.
                                                                                         04-AUG-1995
                                                                                                                  AAQ75585
                                                                                                                                            AAQ75585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 1; Page 10; 39pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-036665/05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (CROS/)
(KURF/)
(BATT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 10-JUN-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W09429723-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   immunodetection; estradiol; alpha-anomer;
                                                                                                                                                                                                                                                                                                                                     (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (PIGA/)
                                                                                                                                                                                                                                 5393
                                                                                                                                                                                                          20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CROS P.
KURFURST R.
BATTAIL N.
PIGA N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Kurfurst R,
                                                                                                                                          standard;
                                                                                                                                                                                                                         AAAAAATACAAAAAAGAAA 5412
                                     gene expression; reverse
                          restriction enzyme; ss.
                                                                                                                                                                                                                                                            Conservative
                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93FR-00007093
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94WO-FR000689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in the alpha-anomer form" 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
                                                                                                                                            DNA;
                                                                                                                                                                                                                                                                      0.3%;
                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           base= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Battail N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "the glycosidic bonds between nucleotides are all
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "carries a
                                                                                                                                            20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         σ
                                                                                                                                            ₽₽
                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                        Score 15.2; DB 1
Pred. No. 9.3e+02
                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Piga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            group derived
                                      transcription; primer; cDNA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ffrom aminopropanediol"
                                                                                                                                                                                                                                                                                    Length
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RESULT 822
AAQ755568/c
ID AAQ755
XX AAQ755
XX AAQ755
XX AAQ755
XX AAALY8
XX AAALY8
KW AAALY8
KW AGGREG
XX JP0630
XX JP0630
XX JP0630
XX JP0630
XX JF0630
XX JF0630
XX JF0630
XX AAALY8
PF 16-APF
XX (NITE
XX (NITE
XX (NITE
XX AAALY8
PT by dig
XX AAALY8
PT D18clc
XX A metl
CC double
CC label:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 밁
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A method for the analysis of cDNA comprises (a) preparing an aggregate of double-stranded cDNAs by using an aggregate of mRNAs and a plural type (labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798) and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates of the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in separate lanes. The
 A method for the analysis of cDNA comprises (a) preparing an aggregate double-stranded cDNAs by using an aggregate of mRNAs and a plural type labelled reverse transcription primers (GENESEQ files AAQ75547-Q75798)
                                                                                                                                                                                                                                                    01-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 5; 11pp; Japanese.
                                                                                        Analysis of cDNA and gene expression - by digestion with restriction enzymes.
                                                                                                                                                                                           16-APR-1993;
                                                                                                                                                                                                                                                                                JP06303997-A.
                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                 04-AUG-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAQ75568;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAQ75568 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 1 A; 0 C; 0 G; 19 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Analysis of cDNA and gene expression -
by digestion with restriction enzymes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-1994
                                                                                                                                  WPI; 1995-018287/03.
                                                                                                                                                                                                                      16-APR-1993;
                                                                                                                                                                                                                                                                                                                                        aggregate;
                                                                                                                                                                                                                                                                                                                                                       Analysis;
                                                                                                                                                                                                                                                                                                                                                                                    Reverse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              method can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NITE ) NIPPON TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JP06303997-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5389 AATTAAAAAATACAAAAA 5408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17;
                                                                                                                                                               NIPPON
                                                                                                                                                                                                                                                                                                                                                                                 transcription primer used in cDNA analysis technique.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AATAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                        gene expression; reverse transcription;
; restriction enzyme; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             be used to analyse gene expression rapidly and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                           Page 5; 11pp;
                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                               TELEGRAPH & TELEPHONE CORP
                                                                                                                                                                                           93JP-00112515.
                                                                                                                                                                                                                      93JP-00112515
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       85.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
                                                           Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.2; DB 1;
Pred. No. 9.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amplification
                                                                                                     amplification of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length
                                                                                                                                                                                                                                                                                                                                                      primer; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of mRNA followed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              easily
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                  유 유
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RESULT 823
AAQ90405/c
ID AAQ904
XX AAQ904
XX AAQ904
XX OB-JAN
XX T2; HL
XW E8:
OS Synthe
XX Synthe
XX Synthe
XX WO9512
XX WO9512
XX WO9512
XX WO9512
XX WO9512
XX WO9512
XX INANO-
XX INAN
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Best Local (
                                                  The sequences represented by, AAQ90402-15 are synthetic DNA probes containing 5' amino termini. The sequences shown in AAQ90390-401 are synthetic DNA probes with 3' ribonucleoside termini. These sequences were specific for the polymorphisms of HLA gene dQa. The sequences were used in the device of the invention. This is a self-addressable electronic device (SAED) that can be used to carry out multi-step and multiplex reactions, such as nucleic acid hybridisations. The advantages of this method are that these reactions can be carried out with complete and precise electronic control, and that the rate, specificity and sensitivity of these reactions are greatly improved at micro-locations
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and using the aggregate of mRNAs as the template for each reverse transcription primer; (b) digesting each of the prepared aggregates the double-stranded cDNAs with restriction enzyme and; (c) electrophoresing the digested aggregate of cDNAs in seperate lanes. method can be used to analyse gene expression rapidly and easily
                                                                                                                                                                                                                                                                                                                                                                                                                           New self-addressable electronic devices - used for multi-step and multiplex reactions such as DNA hybridisation(s), clinical diagnostics and bio:polymer synthesis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 41; 86pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Heller MJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26-OCT-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   11-MAY-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            T2 (synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-JAN-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAQ90405 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NANO-) NANOGEN INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5400 TACAAAAAGAAAAATGAA 5419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1995-185870/24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          dQa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TACAAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          self-addressable electronic device; SAED; hybridisation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA probe with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ĸ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note=
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Pred. No. 9.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino terminal
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Best Local Similarity

0.3%;

Score Pred.

15.2; DB 1; No. 9.3e+02; DB 1;

Length 20;

Query Match

Sequence 20

BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

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RESULT 825
AAV07752/c
ID AAV07752 standard; DNA; 20 BP.
XX
AC AAV07752;
XZ
DT 07-DEC-1998 (first entry)
XX
DS Phosphorothioate oligonucleotide
XX
XX
XX
XX
Phosphorothioate; sulphurisatior
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AAT33632/c
ID AAT336
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Best Local Similarity
Matches 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A PCR primer (AAT33632) is based on rat oncofoetal protein p65 cDNA (see also AAT33630). It was used as the antisense primer together with 5'AmpliFINDER anchor primer (Clontech) to amplify cDNA from the human breast cancer cell line MCP-7. A cDNA clone (AAT33631) coding for human p65 was isolated. p65 is useful as a tumour marker (see also AAW04165).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oncofoetal protein p65; tumour marker; breast cancer; prostate; ovary cancer; diagnosis; prognosis; probe; primer; PCR;
    phosphorothioate; sulphurisation; heterocycle; automated synthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 6 C; 8 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New nucleic acid encoding the p65 oncofoetal protein and related hybridisation probes - for diagnosis of cancer, esp. where relatsteroid hormone expression, and for risk assessment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hanausekwalaszek M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-MAR-1995;
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12-DEC-1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT33632 standard;
                                           Phosphorothioate oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 8; Page 25; 55pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Updated on
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                                                                                                                                                                                                                                                                                            3119 CCCTGACCGAGCTGGACCTG 3138
                                                                                                                                                                                                                                                                   20 CCCGGTCCTAGCTGGACCTG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-MAR-2003 to correct PI field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95US-00405648
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CDNA;
                                                                                                                                                                                                                                                                                                                                                                   0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Slaga TJ,
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                                                                                                                                                                                                                                                                                                                                                                   Score 15.2; DB 1; Length 20; Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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    RESULT 826
AATG3649/c
ID AATG36
XX AATG36
XX AATG36
XX AATG36
XX AATi-H
XX AATi-H
XX AATi-H
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XX human
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention provides a method for sulphurising phosphorus-containing compounds. It comprises contacting the phosphorus-containing compound which a 1,2,4-dithiazolldine-2,5-dione compound or 3-substituted-1,2,4-dithiazolln-5-one compound. The method is especially useful for incorporation of phosphorothioate linkages into biologically important molecules such as DNA, RNA and phosphopptides. Molecules containing such linkages are useful e.g. as antisense compounds for inhibiting gene expression, as reagents for studying DNA-protein or RNA-protein interactions, or as catalytic RNA. The present sequence represents an ollgonucleotide with phosphorothioate linkages prepared by the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sulphurisation of phosphorus-containing oligo:nucleotide(s) - by contacting the containing five-membered heterocycle.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 1 A; 0 C; 0 G; 0 T; 19 U;
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      09-AUG-1995;
                                             25-FEB-1997
                                                                                   JP09052898-A
                                                                                                                          Synthetic.
                                                                                                                                                               antisense; complementary; tax gene; inhibit; HTLV-1;
human T-cell lymphotropic virus type 1; viral antige
                                                                                                                                                                                                                            Anti-HTLV antisense reference oligonucleotide
                                                                                                                                                                                                                                                                   06-JUN-1997
                                                                                                                                                                                                                                                                                                           AAT63649;
                                                                                                                                                                                                                                                                                                                                                  AAT63649 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 7; Page 30; 51pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Barany G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MINU ) UNIV MINNESOTA.

(LOUU ) UNIV LOUISIANA STATE & AGRIC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                             lymphotropic virus
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        95JP-00224606
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1. .20
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                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.3%; Score 15.2; DB 1; 85.0%; Pred. No. 9.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chen L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   compounds, e.g. compound with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0 Other;
                                                                                                                                                                                                                              끍
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hammer RP;
                                                                                                                                                                  antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        рu
                                                                                                                                                                  expression;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        di:sulphide-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <u>,</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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RESULT 827
AAV34591
AAV34591
AC AAV345
AC AAV3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotides having a partial sequence consisting of at least 15 bases of ANT63641 (an antisense oligo complementary to a region of the tax gene which can inhibit human Teal Iymphotropic virus type 1 (HTLV-1) viral antigen expression) are claimed. In an example, six antisense oligos were designed, T1-T6 (ANT63650-55) and were compared to six oligos derived from other regions of HTLV-1, i.e. SJ1 (splice junction), P1 (p21), R1 (rex), RR1 (rex response element), E1 (env) and G1 (gag), four reference oligonucleotides TIS (tax-sense), HC (dC20), HT (dT20) (ANT63647-49) and a random 20mer (RAN) in a HTLV-1 virus antigen expression inhibiting test. Oligonucleotide T1 gave the best results
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mycobacterium vaccae; antigen; therapy; prevention; cytokine production; M. avium; M. tuberculosis; immune response enhancer; cell proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 0, A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                       Mycobacterium vaccae polypeptides - used to develop products for use in detection, therapy and prevention of mycobacteria infections or as immuresponse enhancers.
                                                                                                                                                                                                                                                                                                                                                                             29-AUG-1996;
12-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9808542-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mycobacteria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     M. vaccae antigenic sequence hybridising oligo AD12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAV34591;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 1; Page 8; 10pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anti-HTLV-1 anti-sense oligo:nucleotide - is complementary to region of tax gene from human T-cell lymphotropic virus type 1 and inhibits viral
                                                                                                                                                                                                                                                                Tan P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mycobacterium vaccae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV34591 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          09-AUG-1995;
                                                                                                                                                                                                             WPI; 1998-216926/19
                                                                                                                                                                                                                                                                                                                          (GENB-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOYA-) SOYAKU GIJUTSU KENKYUSHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5393 AAAAAAATACAAAAAAGAAA 5412
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression.
                                                                                                                                                                                                                                                                Hiyama J,
                                                                                                                                                                                                                                                                                                                       GENESIS RES & DEV CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     tuberculosis; immune response enhancer; infection; vaccine; cancer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          95JP-00224606
                                                                                                                                                                                                                                                                                                                                                                                96US-00705347.
97US-00873970.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97WO-NZ000105
                                                                                                                                                                                                                                                                Visser E, Skinner MA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ŗ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                      Scott LM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                   Prestidge RL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0
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                                                                                                                        as immune
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                                                    Mycobacterium vaccae antigens. The invention provides M. vaccae polypeptides that comprise an immunogenic portion of a soluble M. vaccae antigen, or a variant, where the antigen induces an immune response in patients previously exposed to a mycobacterium. Such M. vaccae polypeptides can be used in methods for enhancing non-specific immune response. The methods and products can be used for the detection, treatment and prevention of infectious diseases caused by mycobacteria such as M. vaccae, M. avium or M. tuberculosis. The products also have the ability to induce cell proliferation and cytokine production (e.g. interferon-gamma and interleukin-12 production) in T cells, NK cells, B cells, or macrophages. They can be used for enhancing immune responses for use in vaccines or immunotherapy of infectious diseases and cancers
Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This oligonucleotide is used in the DNA cloning strategies of the
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                Query Match
Best Local S
Matches 17
 5393 ААААААТАСАААААА
                Similarity
                 Conservative
                        0.3%;
                 0,
                         Score
Pred.
                 Mismatches
                         15.2; DB 1;
No. 9.3e+02;
                                 1:
                               Length 20;
                  Indels
                  0
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AAAAAAAAAAAAAAAAAA 20

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Adapter primer oligonucleotide #12 for CAG repeat analysis
                               10-MAR-1999
                                                            AAV68373
                                                                                          AAV68373 standard;
                               (first entry)
                                                                                            DNA;
                                                                                            20
                                                                                            ВÞ
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CAG repeat; human; genome analysis; adapter primer; medical diagnostic; nucleic acid analysis; variation assessment; neurological disease; Huntington's chorea; PCR suppression; ss.

WO9849345-A1

29-APR-1998; 05-NOV-1998 98WO-US008616

29-APR-1997; 97US-0045078P

(UYBO-) UNIV BOSTON

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WPI; 1998-594983/50

Analysing nucleic acid samples - using amplification primers which contain CAG or CTG tri:nucleotide repeats for differential display samples from different sources. of.

Example; Page 31; 44pp; English.

This sequence represents an adapter primer oligonucleotide. It was used to isolate CAG repeat containing sequences from the human genome to test the method of the invention. The method is for analysing nucleic acids in a sample, and comprises: (a) providing a sample containing nucleic acid, a first oligonucleotide primer comprising a CTG repeat, a second oligonucleotide primer comprising a CTG repeat, a second coligonucleotide primer comprising a CTG repeat and a polymerase and PCR reagents; (b) preparing said nucleic acid so that it is amplifiable; (c) amplifying the nucleic acid with the first and second primers; and (d) detecting the amplified product. The method is used to distinguish between the expression of genes in two or more biological samples, e-g. body fluids, cells, solid tissue or solid and liquid foods. It can be used in medical diagnostics, e-g. to differentiate between normal and

Example 8; Page 99; 153pp; English

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RESULT 829
AAT86606/c
ID AAT866
XX AAT866
XX AAT866
XX AAT866
XX Capill
KW Capill
KW Polyac
XX Synthe
PN WO9745
XX WO9745
XX HOVS
PF 23-MA)
XX 24-MA)
XX HOVS
PF 23-MA)
XX Examp:
XX PF A Muscat
XX WPI;
PT Affin:
PT target
XX Separ:
PT target
XX Se
                   CC process using capillary affinity gel electrophoresis. The invention of cleentrically charged target molecules in an analytical mixture. It comprises capillary affinity gel clectrophoresis using a capillary tube which is at least partly filled control to the polymer gel. Receptors for target molecules are covalently bound to the polymer gel. Receptors for target molecules are covalently bound to the polymer. An electric field of at least 50 volts/cm is applied. The capillary tube is charged with the analytical mixture. In a first compliant to the separation stage, the target molecules in the mixture are bound to the compliant to the target molecules are changed, optionally whilst splitting open. In a second stage, the elution conditions are changed, coptionally in stages, so that the affinity of the target molecules for the receptor is eliminated and the target molecules are eluted and control of optionally whilst splitting open. The process is useful for selective separation and/or determination of charged organic compounds, such as oligonucleotides, peptides or carbohydrates. It may be used, e.g. for isolation of specific proteins are carbohydrates. It may be used, e.g. and the carbohydrates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 diseased tissue or to assess the variation within monozygotic twin pairs. The method allows the isolation and analysis of genome subsets containing CAG repeats which are known to be important in a number of neurological diseases including Huntington's chorea. The method uses PCR suppression, in which only fragments which contain a target repeat are efficiently amplified. This allows accurate identification of differentially expressed genes in various cell types. Genome complexity is reduced by the new method which targets genomic subsets containing CAG repeats
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Separation of electrically charged target molecules - by capillary affinity gel electrophoresis using polymer-gel to which receptors for target molecules are bound.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A mixture of oligonucleotides (AAT86604-7) were separated by a new
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example D3; Page 25; 41pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Muscate A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO9745721-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Capillary afinity gel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide separated by capillary affinity gel electrophoresis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-JUN-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1998-041763/04.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-MAY-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (NOVS ) NOVARTIS AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2641 CTGCAGCTGCTGCAGCC 2660
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Paulus A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96CH-00001320
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97WO-EP002647.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
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Pred. No. 9.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 separation; polymer-gel;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length
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Best Local Similarity
Matches 17; Conserv
                                            This invention describes a novel beacon protein and its encoding nucleic acid which is expressed in larger amounts in hypothalamus tissue of obese animals compared to lean animals. Agonists and antagonists of beacon can be used to treat obesity, anorexia, weight maintenance, energy imbalance, diabetes, metabolic syndrome, dyslipidemia, hypertension and/or insulin resistance. The beacon protein, itself is used to manufacture medicaments for treatment of obesity, anorexia, weight maintenance. The treatment is contemplated for both human and animals, such as those important to the livestock industry. The antibody and polynucleotides are useful in diagnosis of conditions a shove. This sequence represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Beacon; hypothalamus; obese; lean; agonist; antagonist; treatment; obesity; anorexia; weight maintenance; energy imbalance; diabetes; metabolic syndrome; dyslipidemia; hypertension; insulin resistance medicament; livestock; diagnosis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                inhibitors. The process achieves higher resolution and selectivity than prior art processes, especially in the case of complex biological prior art processes, especially in sensitivity, even with small amounts of analytical mixtures. It has high sensitivity, even with small amounts of samples. The derivatised polymers may be synthesised specifically using
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAX57365 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard methods
                                                                                                                                                                                                                       Example
                                                                                                                                                                                                                                                     diabetes and metabolic energy levels.
                                                                                                                                                                                                                                                                     New gene encoding a beacon protein associated
                                                                                                                                                                                                                                                                                                      WPI; 1999-337484/28
                                                                                                                                                                                                                                                                                                                                   Zimmet
                                                                                                                                                                                                                                                                                                                                                                               (-ITTI-)
                                                                                                                                                                                                                                                                                                                                                                                                               31-OCT-1997;
11-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              30-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P. obesus beacon PCR primer 3.
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                                                                                                                                                                                                                                                                                                                                                                 (UYDE-) UNIV DEAKIN.
                                primer used in the method of the invention
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                                                                                                                                                                                                                       7; Page 52; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                 INT DIABETES INST.
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97AU-00000323.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.2;
Pred. No. 9.
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                                                                                                                                                                                                                                                                        modulation of obesity,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         resistance
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Query Match Best Local S Matches 17

Similarity

85.0%; 0.3%;

Score 15.2; Pred. No. 9 3 T; 0 U;

.3e+02

Length 20;

0

Mismatches

Indels

0

Gaps

0

Sequence 20

BP; 5

A;

9 C; 3

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0 Other; DB 1;

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RESULT 831
AAX27533/c
ID AAX27533/c
XX AAX275
XX Synthe
XX Silylo
XX Synthe
XX Synthe
XX Synthe
XX Cyanoe
XX Synthe
XX HO9909
XX (PITS/PA (WEIS/PA (JENN/YA)
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                                                                                                                                                                                                                                                                                      The invention relates to silyloxymethyl protected D- or L-ribonucleosides and their phosphonates (I), and silyloxymethyl halides (II). (I) are intermediates for synthesis of RNA-oligonucleotides with predetermined nucleotide sequence, particularly by machine synthesis. The groups specified above, apart from those on silyl, are those particularly for the cyanoethyl phosphoramidate coupling. Uses of the oligoribonucleotide products in diagnosis, therapy, and as research tools, are well known, and are mot dealt with in detail. (II) is an intermediate for (I). The silyloxymethyl halide reagent is easy to prepare, and yields are high. Introduction of the silyloxymethyl group into the ribonucleoside is simple and rapid, and the acetal bond formed does not migrate, eliminating particularly the prior art problem of 2' to 3' isomerisation. The methylenedioxy group spacer between the silyl group and nucleoside ring results in less steric hindrance than bulky direct silyloxy to provide, e.g., acid or base stability, and second, higher yields in coupling. Purer products are therefore obtained than in prior art, canabling larger quantities and longer chains of oligoribonucleotides to be synthesised successfully, and in shorter times
                                                                                                                   Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2-Silyloxymethyl ribonucleosides and their phosphonate derivatives - have high purity, use in machine synthesis of ribonucleic acids, enable longer oligonucleotide chain construction, and larger amounts.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Silyloxymethyl; cyanoethyl phosp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic RNA sequence produced by
                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 6; Page 25; 38pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-180963/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pitsch S, Weiss PA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (PITS/) PITSCH S. (WEIS/) WEISS P A. (JENN/) JENNY L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 17-AUG-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18-AUG-1997;
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                                                         5393 AAAAAAATACAAAAAAGAAA 5412
                                                                                                                                                   Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          thyl; phosphonate; silyloxymethyl halide; diagnosis; ss; phosphoramidate coupling; isomerisation; steric hindrance.
                                                                                                                                                                                                                                         BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                         0 A;
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                                                                                                                                            0.3%;
                                                                                                                                                                                                                                         0 C; 0 G; 0 T;
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                                                                                                                                                Score 15.2;
Pred. No. 9.
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                                                                                                                      Mismatches
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                                                                                                                                                                                                                                         20 U; 0 Other;
                                                                                                                                                   9.3e+02;
                                                                                                                                                                            DB 1; Length 20;
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RESULT 832
AAZO4702/c
ID AAZO47/
XX PCR pr
XX Vaccin
XW PCR pr
XX Vaccin
XW Dartho
XX Synthe
OS Synthe
OS Synthe
OS Synthe
OS Synthe
OS Synthe
OS Synthe
OX (CLlamy)
XX WO9928-
XX I1-JUN
XX II-JUN
X
RESULT 833
AXX27889
ID AAX278
XX
AC AAX278
AC AAX278
AC O2-JUN
XX
DE Probe
XX
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17-DEC-1997;
04-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCR primers AAZ01426-Z06209 were used to amplify open reading frames (ORFS) of the genome of Chlamydia trachomatis (see AAZ01425). These ORFS encode polypeptides (see AAY36754-Y37949) which can be used as vaccines against Chlamydia trachomatis. Antisense and ribozyme sequences can also be used to control growth of the microorganism. Chlamydia trachomatis is responsible for a large number of diseases, e.g. eye diseases such as conventional trachoma, nonendemic trachoma, paratrachoma, and inclusion conjunctivitis; genital diseases such as nongonococcal uretritis, pepidymitis, cervicitis, salpingitis, perihepatitis, bartholinitis; pneumopathy in breast feeding infants; and venereal lymphogranulomatosis.
                                                                                          02-JUN-1999
                                                                                                                                                                                                          AAX27889 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 7 A; 3 C; 7 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 1710; 1755pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Genome sequence of Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1999-371125/31.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer used to amplify an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07-OCT-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9928475-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bartholinitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  paratrachoma; inclusion conjunctivitis; genital disease; perihepatitis;
nongonococcal uretritis; epidymitis; cervicitis; salpingitis; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                              5326 GCAGGCTTTCCAGTTTTTAC 5345
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17; Conserv
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                                                                                       (first entry)
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97FR-00016034.
98US-0107077P.
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                                                                                                                                                                                                          ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Mismatches
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Pred. No. 9.
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Probe for human

CSR

protein coding

sequence.

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RESULT 834
AAZ11326
ID AAZ113
XX AAZ113
XX AZ113
XX Mycob
DT 25-OCT
XX Mycob
KW Mycob
KW Mycob
KW Iespi
KW Iespi
KW 1espi
KW 1esp
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This sequence represents a probe for DNA encoding a human cellular stress response (CSR) protein of the invention. The CSR proteins are macrophage scavenger receptor proteins. The CSR proteins can be used in the treatment, gene therapy and diagnosis of diseases in which intracellular stress is important, such as arteriosclerosis, diabetic circulatory obstruction, and microbial infection. Expression of the proteins is induced in vivo in response to intracellular stress, and inhibits cell death as a result of such stress
                                                                                                                                                                                                                                                                                                                            dendritic cell maturation; infectious disease; immune disorder; cancer; respiratory system; mycobacterial infection; allergy; tuberculosis; leprosy; sarcoidosis; lung cancer; asthma; skin disorder; psoriasis; dermatitis; eczema; alopecia areata; skin cancer; basal carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 8 A; 3 C; 6 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Scavenger receptor proteins - for treatment and diagnosis of disorders involving cell stress.
                                       23-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mycobacterium vaccae protein; antigen; T cell activation; cytokine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mycobacterial 16S rRNA specific oligo AD12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 10; Page 167; 175pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nakamura Y,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9909159-A1
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                                                                                              01-JUL-1999.
                                                                                                                                                        WO9932634-A2
                                                                                                                                                                                                                 Mycobacterium vaccae
                                                                                                                                                                                                                                             Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAZ11326 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
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                                                                                                                                                                                                                                                                                                      squamous cell
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NISB)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    568 CTGAAGAAGGAGGAGCTGAA 587
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
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                                                                                                                                                                                                                                                                                                      carcinoma;
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98JP-00230121.
                                       98WO-NZ000189.
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85.0%;
                                                                                                                                                                                                                                                                                                      melanoma;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.2;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                   PCR primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    .3e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length
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RESULT 835
AAV64851
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XX AAV648
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23-DEC-1997;
23-DEC-1997;
11-JUN-1998;
17-SEP-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pharmaceutical compositions comprising the proteins or nucleic acid sequences encoding the proteins can be used for the treatment, prevention, and detection of disorders including infectious diseases, immune disorders and cancer. In particular, the compounds and methods used for treatment of diseases of the respiratory system, such as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B. cereus 16S rRNA oligonucleotide primer #8.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mycobacterial infections, asthma, allergies, tuberculosis, leprosy sarcoidosis and lung cancers, and disorders of the skin such as psoriasis, atopic dermatitis, eczema, allergic contact dermatitis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cells and natural killer cells, to stimulate the production of cytokin to enhance the expression of costimulatory molecules on dendritic cel and monocytes, and to enhance dendritic cell maturation and function proteins can be expressed by standard recombinant methodology.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Tan
                                                      WPI; 1999-038276/04.
                                                                                                                                                         30-APR-1997;
                                                                                                                                                                                                       30-APR-1997;
                                                                                                                                                                                                                                                                                                       JP10295377-A.
                                                                                                                                                                                                                                                                                                                                                         Bacillus cereus.
                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV64851;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 carcinoma and melanoma
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alopecia areata, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention provides heat-killed Mycobacterium vaccae, or recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 15; Page 177; 243pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Enhancing
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                                                                                                     (QPPP ) QP CORP
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nes 17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                    primer;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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97US-00997080.
97US-00997362.
98US-00095855.
98US-00156181.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         and skin cancers such as basal carcinoma, squamous
                                                                                                                                                                                                                                                                                                                                                                                                                                 detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Visser ES,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; 0 C; 0
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Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                    PCR; microbe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             an antigen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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An oligonucleotide primer - useful for detection of Cereus microbes.

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RESULT 836
ANX21950/c
ID ANX219
XX ANX219
XX ANX219
XX Human
XX Human
XX Antis
KW An
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the presence of
detect Bacillus
                      This sequence represents an example of an antisense oligonucleotide of the invention. The oligonucleotides are 8-50 nucleotides in length, and are targeted to a nucleic acid encoding human B-raf and which is capable of inhibiting human B-raf expression. The oligonucleotides is used to inhibit the (abnormal) expression of human B-raf, to inhibit hyperproliferation of cells, to treat or prevent an abnormal proliferative condition, e.g. hyperproliferative disorders such as cancer (e.g. of the brain or nervous system), restenosis, psoriasis or a disorder characterised by T-cell activation and growth. They may also be used to diagnose these diseases, as well as atherosclerosis. The oligonucleotides of the invention may be used to distinguish raf-associated tumours from tumours having other etiologies. The antisense oligonucleotides can also be used to quantify raf expression in assays
                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New antisense oligonucleotides - for modulation of human B-raf
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1999-120502/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antisense oligonucleotide; B-raf; human; inhibitor; T-cell activation; hyperproliferative disorder; cancer; restenosis; psoriasis;
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                                                                                                                      Electrochemical detection; glucose; cholesterol; urea nitrogen; bilirubin; uric acid; haemoglobin; lactic acid; body fluid; blood; plasma; serum; urine; lymph diagnosis; ss.
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                                                                                                                                                                                                                                      detection method sample DNA target.
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98US-0107078P
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RESULT 840
AAA40448/C
ID AAA404
AC AAA604
AC AAA604
AC AAA604
XX 13-NOV
XX Electr
XW Electr
XW biliru
XW b
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for quantitatively analysing a blochemical compound (1) which comprises contacting (1) with double constrained DNA fixed to the surface of an electrode at their terminals in the combine comprises contacting (1) with double combined to the surface of an electrode at their terminals in the contact of the compound and becomes reduced, and detecting electric potential to the compound and becomes reduced, and detecting electric current flowing the between the electrode and a second electrode in the aqueous medium. The combined is useful for detection of blochemical compounds such as glucose, coholesterol, urea nitrogen, blirubin, uric acid, haemoglobin and lactic acid in body fluids such as whole blood, plasma, serum, urine, and lymph for diagnosis of various diseases. The method allows detection of blochemical compounds quickly and easily with a high sensitivity using a simple apparatus. This sequence represents DNA fragment used as a target of the compounds of the invention
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                          Electrochemical detection; glucose; cholesterol; urea nitrogen; bilirubin; uric acid; haemoglobin; lactic acid; body fluid; blood;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20
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                                   06-JAN-1999;
24-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Electochemical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA40448 standard;
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                                                                                                                         07-JAN-2000; 2000EP-00100126.
                                                                                                                                                                                                                                                                                                                                                                          plasma; serum;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; Page 8; 14pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (FUJF ) FUJI PHOTO FILM CO LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000-444372/39.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                   99JP-00001111.
99JP-00143599.
                                                                                                                                                                                                                                                                                                                                                                             urine; lymph diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    detection method
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA;
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Pred. No. 9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    fixed probe
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RESULT 841
ABL41405
ID ABL414
XX ABL414
XX ABL414
XX Primer
XX VIOLENT
DE Univer
XX VIOLENT
DE Unider
XX VIOLENT
XX VIOLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes a novel method for quantitatively analysing a blochemical compound (I) which comprises contacting (I) with double stranded DNA fixed to the surface of an electrode at their terminals in which electrochemically active threading intercalators are intercalated, in an aqueous medium under application of electric potential to the electrode in the presence of an oxidase which oxidizes the blochemical compound and becomes reduced, and detecting electric current flowing between the electrode and a second electrode in the aqueous medium. The method is useful for detection of blochemical compounds such as glucose, cholesterol, urea nitrogen, bilirubin, uric acid, haemoglobin and lactic acid in body fluids such as whole blood, plasma, serum, urine, and lymph for diagnosis of various diseases. The method allows detection of blochemical compounds quickly and easily with a high sensitivity using a simple apparatus. This sequence represents DNA fragment used as fixed probe DNA in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                         02-MAY-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                02-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Primer; universal
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                                                          Disclosure, Page 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    KR98082207-A
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                                                                                                                    Universal primer
                                                                                                                                                                                 WPI; 2000-069105/06
                                                                                                                                                                                                                                        Kang HW,
                                                                                                                                                                                                                                                                                                  (RURA-)
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                                                                                                                                                                                                                                                                                               RURAL DEV ADMINISTRATION.
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                                                                                                                                                                                                                                                                                                                                                         97KR-00016981.
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                                                                                                                                                                                                                                        Eun
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for the genetic diagnosis of species
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
                                                       21pp; Korean.
                                                                                                                 genetic diagnosis
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                                                                                                                                                                                                                                        Koh SJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                       유
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The invention relates to universal primers for the genetic diagnosis

also be used in kits

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ARACIDAT 842
ARALIZO81/c
ID ARALIZO
XX ARALIZO
XX ARTISE
XW WOZOOO
XX OG-APR
PP 21-SEP
PR 04-DEC
PR 08-JUN
XW WPI; 2
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                                                                                  This invention describes novel antisense nucleic acids (I) targeted (C) against a specific nucleic acid sequence within human ICAM-1. The CC products of the invention have cytostatic, anti-inflammatory, CC dermatological, antiviral, antirheumatic, antiarthritic, CC dermatological, antipsoriatic, antiasthmatic, antitussive activity and CC can be used for antisense therapy; gene therapy. The antisense nucleic CC acids or vectors are used to inhibit or eliminate acute or chronic CC inflammatory diseases of humans, virus infection, metastasis, inflammation of the skin, mobilization of hematopoietic stem cells, CC coughs and all biological process under the influence of ICAM-1, CC ulcerative colitis, rheumatoid arthritis, lupus erythematosus, organ CC rejection, graft-versus-host reaction after bone marrow transplantation, CC psoriasis, asthma and neurodermatitis. In particular, the antisense concleic acids are used to treat acute or chronic inflammation of gum disease. The antisense nucleic acids regulate or suppress ICAM-1 gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 antitussive gene therapy; inflammatory disease; virus infection; metastasis; hematopoietic stem mobilization; ulcerative colitis; rheumatoid arthritis; lupus erythematosus; organ rejection; psor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    species. The current sequence represents a universal primer for the genetic diagnosis of species
                                  expression. The antisense nucleic acids or vectors can be used to suppress immune reactions against gene therapy using viral or non-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antisense nucleic ICAM-1 gene, useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-JUN-1999;
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04-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antisense; primer; ICAM-1; human; adhesion molecule; cytostatic; anti-inflammatory; dermatological; antiviral; antinheumatic; asthma; antisarthritic; immunosuppresive; antipsoriatic; antiasthmatic; antitussive gene therapy; inflammatory disease; virus infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human ICAM-1 antisense oligonucleotide 650D.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene, useful
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 BP; 2 A; 3 C; 9 G; 6 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24; 28pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98DE-01044111.
98DE-01056138.
99DE-01026110.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           acids targeted to specific sequences within the treating inflammation and metastasis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Steidl U,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ВP
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Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 U;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .3e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 gum disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sczakiel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             rejection; psoriasis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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RESULT 844
AAC78948/c
ID AAC78948
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AAZ37992/c
ID AAZ379
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                                                                                                           Query Match
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Matches 17
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Glaucoma;
GLC1A gene
                                                                                                                                                                               The invention relates to a method for the determination of a predisposition to glaucoma. The method comprises amplifying a GLC1A gene with a primer pair selected from the sequences shown in AAZ37981-Z38008. The primers are used to determine whether a subject has or has the potential to develop primary open wide angle glaucoma. Sequences AAZ37981-Z38008 represent primer pairs specific for human GLC1A gene exon sequences. These primers were used for the GLC1A assay to identify patients having a predisposition to glaucoma
                                                                                                                                                           Sequence 20 BP; 4 A; 8 C; 4 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20
                                                                                                                                                                                                                                                                                             Claim 1; Page 131; 137pp; English.
                                                                                                                                                                                                                                                                                                                               Determination of a predisposition to
                                                                                                                                                                                                                                                                                                                                                                               Stone EM,
                                                                                                                                                                                                                                                                                                                                                                                                                               07-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9951779-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human GLC1A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-FEB-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAZ37992 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   to diagnose ICAM-1 associated disturbances. AAA12063-A12092 and AAA12099 represent the antisense oligonucleotides described in the method of the
                                                                                                                                                                                                                                                                                                                                                                                                        (IOWA ) UNIV IOWA RES FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
les 17; Conserv
                                                                           1960 GGGTTCTCTGAGTCCAGCAG 1979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2844 GCACAGATCAACATGGAGCC 2863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ma; PCR amplification; primary open wide angle glaucoma;
gene; human; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
                                                             20
                                                                                                             17;
                                                                                                                       Similarity
   standard;
                                                             GGGGACTCTGAGTTCAGCAG 1
                                                                                                                                                                                                                                                                                                                                                                               Sheffield VC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 2
                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                               98US-00056285
                                                                                                                                                                                                                                                                                                                                                                                                                                                       99WO-US007671.
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  DNA;
                                                                                                                      0.3%;
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  ВÞ
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                                                                                                                      Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 15.2; DB 1
Pred. No. 9.3e+02
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                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                               WLM,
                                                                                                                                                                                                                                                                                                                               glaucoma by analysing mutations
                                                                                                                                                                                                                                                                                                                                                                               Fingert
                                                                                                                                   DB 1; Length 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
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5196 TCAGCGTGGGAGGCCACGTG 5215

Query Match Best Local S Matches 17

l Similarity 17; Conserv

Conservative

0;

Mismatches

0.3%;

Score 15.2; DB 1; Pred. No. 9.3e+02;

Length Indels

20;

0

Gaps

0,

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Ashkenazi AJ, Baker KP,
Ferrara N, Filvaroff E,
Goddard A, Godowski PJ,
Kljavin IJ, Kuo SS, Napi
Stewart TA, Tumas D, Wil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        28-APR-1999;
14-MAY-1999;
23-JUN-1999;
26-JUL-1999;
29-OCT-1999;
30-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                           30-DEC-1999; 99WU-UGUUZ19.
05-JAN-2000; 2000WO-US000219.
06-JAN-2000; 2000WO-US000277.
06-JAN-2000; 2000WO-US000376.
                                       AAC78458 to AAC78599 represent polynucleotide and EST (expressed sequence tag) sequences which encode secreted or transmembrane PRO polypeptides. The PRO polynucleotides and polypeptides have cytostatic activity. The polynucleotides and polypeptides can be used for detecting the presence of PRO polypeptides in samples, for linking bioactive molecules to cells and for modulating biological activities of cells, using the polypeptides for specific targeting. The polypeptide targeting can be used to kill the target cells, e.g. for the treatment of cancers. The polypeptide pairs provide specific targeting of bioactive molecules to cells. AAC78600 to AAC78987 represent PCR primers and probes used in the isolation of the
                                                                                                                                                                                                                                                Novel PRO polypeptides and polynucleotides used in detection methods, to target bioactive molecules to specific cells, and to modulate cellular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-MAR-1999;
12-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human
 Sequence
                                                                                                                                                                                                       Example 114; Page 342; 636pp; English
                                                                                                                                                                                                                                                                                            WPI; 2000-611443/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-FEB-2000; 2000WO-US004341.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expressed
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                                                                                                                                                                                                                                                                                                                                                                                                             (GETH ) GENENTECH INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 16-DEC-1999;
                              polynucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRO772 reverse PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    secreted protein; transmembrane protein; PRO; EST; cytomed sequence tag; detection; cancer; PCR primer; probe;
   20
BP; 4 A; 7 C; 4 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       99US-0145698P.
99US-0162506P.
99WO-US028313.
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99US-0134287P.
99US-0141037P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           99WO-US028551.
99WO-US028565.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 99WO-US030095.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99WO-US031243.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-US005028
                              sequences
                                                                                                                                                                                                                                                                                                                     KR, Botstein D, Desnoyers L, Eaton DL;
f E, Fong S, Gao W, Gerber H, Gerritsen ME;
PJ, Grimaldi CJ, Gurney AL, Hillan KJ;
Napier MA, Pan J, Paoni NF, Roy MA, Shelto,
Williams PM, Wood WI;
 5 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID
   0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NO:577.
                                                                                                                                                                                                                                                                                                                                        Shelton DL;
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Page 413
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RESULT 845
AAZ91117/c
ID AAZ91117;
XX
AC AAZ91117;
XC AAZ91117;
XC AAZ91117;
XC AAZ91117;
XC AAZ91117;
XC AAZ91117;
XX Oligonucleotide #5 for conjugati
XX Abietane derivative; labelling;
XX Abietane derivative; labelling;
XX Abietane derivative; labelling;
XX Abietane derivative; labelling;
XX O4-FEB-2000.
XX Synthetic.
XX Synthetic.
XX S1-JUL-1998; 98FR-00010084.
PF 31-JUL-1998; 98FR-00010084.
PF 31-JUL-1998; 98FR-00010084.
XX INMR) BIO MERIEUX.
XX X
XX (INMR) BIO MERIEUX.
XX X
XX (INMR) BIO MERIEUX.
XX X
XX (INMR) BIO MERIEUX.
XX Saturated and unsaturated derivention privatives with naturated deriventions
XX Saturated and unsaturated deriventions
XX Saturated and unsaturated deriventions
XX Example 5; Page 20; 39pp; Frenc.
XX Charles MH, Piga N, Battail Pl
XX WPI; 2000-239603/21.
XX Saturated and unsaturated derivention relates to novel derivatives. The new compounds of the convention relates to novel derivatives. The object of the produce monoclonal antibodie of the produce monoclonal antibodie of sequences that can be labele conventions with biological mc XX Sequence 20 BP; 0 A; 0 C; 0 G;
SQ Sequence 20 BP; 0 A; 0 C; 0 G;
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                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                          The invention relates to novel saturated and unsaturated abletane derivatives. The new compounds may be used directly or indirectly in the development of new disgnostic tests, to follow infections, especially viral infections, to follow and/or measure chemical products, especially potential pollutants. In diagnostic tests they may be used as markers, or to form a universal solid phase after immobilization on a solid support, to produce monoclomal antibodies or polyclonal antibodies having diagnostic uses. The oligonucleotides AAZ91113-Z91117 represent examples of sequences that can be labeled with the new abletane derivatives. The new derivatives may be used to substitute for biotin in diagnostic tests, but because they are not found naturally in humans the risk of potential interactions with biological molecules is eliminated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saturated and unsaturated derivatives of abjetic acid and their conjugated derivatives with natural and synthetic polymers, having use diagnostics, chemical reactions and analysis.
   Wolframin; human; Wolfram Syndrome;
                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 5; Page 20; 39pp; French.
                                              Human wolframin
                                                                            26-MAY-2000
                                                                                                                                       AAZ88913 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Abietane derivative; labelling; diagnostic test; biotin substitute;
                                                                                                                                                                                                                                          5393 ААААААТАСААААААСАА 5412
                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
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                                                                                                                                                                                                                                                                          Conservative
                                                                            (first entry)
                                              exon
 transmembrane protein; dia psychiatric disease; gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         for conjugation to abietane derivative.
                                                                                                                                                                                                                                                                                      0.3%;
                                             8-3 PCR primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Battail PN,
                                                                                                                                                                                                                                                                        0,
                                                                                                                                                                                                                                                                        Score 15.2; DB 1,
Pred. No. 9.3e+02,
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Veron L,
                                                #2.
dlagnosis; therapy;
ene therapy; primer;
                                                                                                                                                                                                                                                                                                    DB 1; Length 20;
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                                                                                                                                                                                                                                                                          Indels
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                                                                                                                                                                                                                                                                        Gaps
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Best Local (
      WPI; 2000-558188/51.
                                                                                                                                                     17-AUG-2000
                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                07-NOV-2000
                                                                                                                                                                                                                                                                                                                                                                                                              AAA50193;
                                                                                                                                                                                                                                                                                                                                                                                                                                          AAA50193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Strom T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17;
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This invention describes a novel human transmembrane protein, Wolframin. Wolframin and DNA encoding it are useful for diagnosis and/or therapy of Wolfram Syndrome, in particular where there is a tendency to psychiatric disease. The products of the invention can be used for gene therapy. This sequence represents a PCR primer used in the amplification of the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Wolframin, a transmembrane protein and related DNA useful for diagnosis and therapy of Wolfram Syndrome, especially where there is a tendency to psychiatric disease.
Manoharan M, Maier MA;
                                                                                   12-FEB-1999;
                                                                                                                           11-FEB-2000; 2000WO-US003543
                                                                                                                                                                                                                                                                                            modified_base
                                                                                                                                                                                                                                                                                                                                                                                                        Phosphodiester oligonucleotide; H-phosphonate chemistry;
                                                                                                                                                                                                                                                                                                                                                                                                                                                2'-Methoxyethoxy-modified oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 9 A; 1 C; 10 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure, Page 8; 26pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-184136/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
                                                                                                                                                                                                               WO200047593-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 wolframin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYMU-) UNIV MUENCHEN MAXIMILIANS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5121 GCGCAAAGAGGAATAGGAGG 5140
                                         ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GGGCAAAGAGGAAGAAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                            PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                      9908-00250075
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                                                                                                                                                                                                                                                                             /*tag=
                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                       /*tag= a
/note= "2'-methoxyethoxy modified thymidine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
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85.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.2; DB 1
Pred. No. 9.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                           88
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RESULT 848
AAK99515
ID AAK995
XX AAK995
XX URP ol
XX Polyme
XX Polyme
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XX Uni
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CC compounds and methods for the preparation of mixed backbone oligomeric, compounds having phosphodiester internucleoside linkages in CC addition to phosphorothioate and/or phosphoramidate internucleoside linkages. The methods also include incorporation of boranophosphate CC linkages. The methods also include incorporation of boranophosphate are coupled together forming contiguous regions of lor more H-CC phosphonate internucleoside linkages. Bach contiguous region is CC subsequently oxidized to phosphodiester, phosphorothioate, phosphoroamidate or boranophosphate internucleoside linkages prior to CC further elongation. Mixed backbone oligomeric compounds are prepared in CC contiguous region of the invention are prepared using novel oxidation steps that oxidize a region of lor more H-phosphonate internucleoside CC linkages without degrading existing linkages that have been previously coxidized. The oligonucleotides obtained are useful as primers in PCR, CC probes, linkers, gene fragments and for other diagnostic tests on e.g. CC biological tissue, fluid, cells etc., as research reagents, and as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 17
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                               Disclosure; Page 4; 10pp; Korean
                                                                                                                                                        WPI; 2000-609364/58
                                                                                                                                                                                                                                                                                                                                          07-APR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                   07-APR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Polymerase chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           URP oligonucleotide relating to a primer of plasmid subspecies 2F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          06-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAK99515;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             antiviral agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence is that of a phosphodiester oligonuclectide containing 20 T nucleobases, 10 having a 2' methoxyethoxy group on its ribosyl sugar moiety. It is an example of an oligomeric compound production of the methods of the invention. The invention provides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Preparation of mixed backbone oligomeric compounds useful for diagnostic tests, involves oxidation of H-phosphonate linkages to phosphodiester internucleoside linkages.
                                                                                            Polymerase chain reaction (PCR) primer of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 12; Page 34; 49pp; English.
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                                                                                                                                                                                                                                                                                 RURAL DEV
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                                                                                                                                                                                                                 SJ,
                                                                                                                                                                                                                                                                                                                                             98KR-00012278
                                                                                                                                                                                                                                                                                                                                                                                                      98KR-00012278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              reaction; PCR; primer; plasmid subspecies 2F; URP;
                                                                                                                                                                                                                                                                                 ADMINISTRATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                    Kwon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
                                                                                                                                                                                                                    WS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ВP
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Pred. No. 9.3e+02.
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                                                                                            plasmid
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                                                                                            subsp.
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RESULT 849
AAC58208/c
ID AAC582
XX AAC582
XX AC582
XX AC582
XX Luman
DE Human
XX Human;
KW Lumouz
KW Cytost
KW inmunc
XX Homo s
XX WO2000
XX 14-SEE
PN WO2000
XX 16-JAN
XX 08-MAJ
PR 08-MAJ
PR 12-MAJ
PR 12-MAJ
PR 29-MAJ
PR 10-OC1
PR 30-DE(
PR 30-DE
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-APR-1999;
05-OCT-1999;
30-NOV-1999;
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                  The present invention describes an isolated antibody (Ab) that binds to one of the human proteins (P) designated PRO213, PRO1330, PRO1449, PRO237, PRO324, PRO351, PRO351, PRO351, PRO351, PRO351, PRO351, PRO351, PRO538, PRO561 PRO772, PRO772 or PRO474. The Ab can be used in compositions a methods for the diagnosis and treatment of neoplastic cell growth and proliferation in meammals, including humans. Genes and polypeptides encoded by them, that are amplified in the genome of a tumour cell, ce be identified and are useful targets for the treatment and prevention be identified and are useful targets for the treatment and prevention
                                                                                                                                                                                                                                                                                                            New isolated antibody for use in compositions and methods for the diagnosis and treatment of neoplastic cell growth and proliferation mammals, including humans, and in monitoring tumor treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20
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21-APR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; tumour; diagnosis; neoplastic disease; identification; cancer; tumourigenesis; detection; neoplastic cell growth; proliferation; cytostatic; antiinflammatory; immunomodulatory; inflammatory disorder immunological disorder; hybridisation; probe; PCR primer; ss.
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                                                                                                                                                                                                                                                            Example 14;
                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-572269/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Baker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-DEC-1999
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17; Conserv
                                                                                                                                                                                                                                                                                                         including humans,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Desauvage FJ,
                                                                                                                                                                                                                                                          Page 118; 195pp; English.
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99US-0123957P.
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99WO-US031274.
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99WO-US023089.
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Pred. No. 9.3e+02;
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                                                                                                                                                                   The present sequence is that of primer betaseq2, which was used with primer betarev (see AAA99972) in the PCR amplification of a 416 bp fragment of mouse vascular-endothelial protein tyrosine phosphatase (VE-PTP) cDNA (see AAA88865). PCR analysis was used to examine VE-PTP expression in mouse tissues and during mouse embryonic development. In adult mouse, VE-PTP was strongly expressed in brain as well as in lung and heart. In embryonic development, VE-PTP increased from day E11 to day E17. VE-PTP polypeptides, nucleic acids and ligands are used in claimed methods for detecting and modulating receptor tyrosine kinase Tie-2 activity. This allows the monitoring or modulation of anglogenesis, induction or inhibition of vascular growth or remodelling and blood respectively.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Vascular-endothelial protein tyrosine phosphatase; VE-PTP; mouse; Tie-receptor-type tyrosine kinase; antiangiogenic; antitumour; antimatastatic; tumour; metastasis; anglogenesis; therapy; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              can be identified and used as antagonists. Benign or malignant tumours inflammatory disorders and immunological disorders can be treated. AAC58123 to AAC58224 represent hybridisation probes and PCR primers us in the isolation of the human PRO sequences. AAC58225 to AAC58241 and AAB24041 to AAB24056 represent human PRO polynucleotide and protein
                                                                                        Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Monitoring or modulating regulating tumor growth,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-648932/63.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 2; Page 4; 60pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             phosphatase.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PLAC ) MAX PLANCK GES FOERDERUNG WISSENSCHAFTEN.
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                                                                                                                                                    vessel maturation,
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                                                                                        11 C; 2 G; 6 T; 0 U; 0 Other;
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Score 15.2; DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   tyrosine kinase activity, useful e. vascular-endothelial protein tyrosi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                adrenergic receptor, or its W64R mutant, in a medium containing that stimulate differentiation of the cells. The adipocytes are models for studying biological mechanisms in adipocytes, and to for agents potentially useful in treatment of obesity, diabetes hyperlipidemia, specifically antilipid agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PCR primers AAA57407-08 were used to amplify DNA encoding beta-actin. Treactions was performed to analyse and compare the products of adipose tissue. The specification describes a method for the production of adipocytes from non-differentiated fibroblasts. The method comprises culturing confluent fibroblasts that tably express the human beta3 adrenergic receptor, or its W64R mutant, in a medium containing agents that stimulate differentiation of the cells. The adipocytes are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Beta-actin; adipose tissue; adipocyte; fibroblast; beta3 adrenergic receptor; cell differentiation; chyperlipidemia; antilipid; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 3 A; 3 C; 9 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              potential anti-obesity agents, lexpressing the beta3 adrenergic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Production of adipocytes from fibroblasts, useful e.g. for a potential anti-obesity agents, by culturing confluent cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   obesity; diabetes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                screen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The
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01-NOV-2001 AAD14829; AAD14829

(first entry)

standard;

DNA;

20

BP

Human

glycogen

synthase kinase 3 alpha antisense oligo

SISI

#116670

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The invention relates to an antisense compound 8 to 30 nucleobases in length targetted to a nucleic acid encoding glycogen synthase kinase 3 alpha. The antisense compound specifically hybridises with and inhibits
                                                 Antisense compound 8 to 30 nucleobases in length comprising a that is targeted to a nucleic acid molecule encoding glycogen kinase 3 alpha, useful for the treatment of e.g. diabetes and hyperproliferative disorders.
                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            neurological disorder; tumour; haematopoietic disord hyperproliferative disorder; developmental disorder; phosphorothioate backbone; ss.
                                  Example
                                                                                                                                            21-JAN-2000; 2000US-00488856
                                                                                                                                                            16-JAN-2001; 2001WO-US001411
                                                                                                                                                                              26-JUL-2001.
                                                                                                                                                                                             WO200152865-A1.
                                                                                                                                                                                                                             modified_base
                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                      modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; glycogen
                                                                                           2001-442247/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             sapiens.
                                                                                                            BP,
                                                                                                                             SISI
                                 15; Page 84;
                                                                                                          Mckay R,
                                                                                                                             PHARM
                                                                                                                                                                                                                                       /*tag=
'--od_base=
"Me'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             synthase
                                                                                                                                                                                                              /*tag=
/mod_ba
                                                                                                                                                                                                                               /note=
18
                                                                                                                                                                                                                                                                      /*tag=
                                                                                                                                                                                                                                                                                         /mod
                                                                                                                                                                                                                                                                                                                  /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note=
                                                                                                                                                                                                                                                                                                                                                /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mod
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           _bom_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                         _
mod
                                                                                                                                                                                                                                                                                                                                                                                                                         mod_base=
                                                                                                                                                                                                                                                                                                        *tag=
                                                                                                                                                                                                                                                                                                                                                                                                 mod_base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                   note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diabetes; hyperproliferative disorder;
                                                                                                                                                                                                            _base=
                                                                                                                                                                                                                                                                       base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            base= OTHER
                                115pp;
                                                                                                            Butler MM,
                                                                                                                                                                                                                                                                                                base=
                                                                                                                                                                                                                                                                                                                                                                         base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                          раве=
                                                                                                                                                                                                                                                                                                                        base=
                                                                                                                                                                                                                                                                                                                                                 _base=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Phosphorothioate backbone"
                                                                                                                                                                                                                                       "Methoxyethyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Methoxyethyl residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     þ
                                                                                                                                                                                                                                                                                                                                                        ω
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             kinase 3 alpha; antidiabetic; cytostatic;
                                                                                                                                                                                                                                                                                                m5c
                                                                                                                                                                                                              m5c
                                                                                                                                                                                                                                               OTHER
                                                                                                                                                                                                                                                                                                                                                                                                  m5c
                                                                                                                                                                                                                                                                        m5c
                                                                                                                                                                                                                                                                                                                         m5c
                                                                                                                                                                                                                                                                                                                                                 m5c
                                                                                                                                                                                                                                                                                                                                                                         m5c
                                                                                                                                                                                                                                                                                                                                                                                                                          m5c
                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER
                                  English.
                                                                                                           Wyatt JR;
                                                                                                                                                                                                                                       residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     inflammation;
                                                                  synthase
                                                                           compound
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RESULT 853
AAK95023
ID AAK950
XX AAK950
XX AAK950
XX O6-NOV
XX Human
XX Human
XX Human
XX Homo g
XX EP1130
XX 05-SEE
XX 05-SEE
XX 07-JUI
XX 08-JUI
PR 11-JAA
XX 02-MAI
XX WPI;
I Ota T,
PI Wakama
XX WPI;
I Ota T,
PI Sand ni
CC for ci
CC for ci
CC for ci
CC enable
CC enable
CC enable
CC enable
CC clone
XX Seque
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                           08-JUL-1999;
11-JAN-2000;
02-MAY-2000;
                                           The invention relates to primers for synthesising full length cDNA clones, 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special methods. The present sequence is a primer used to amplify a human cDNA clone present sequence is a primer used to amplify a human cDNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the expression of glycogen synthase kinase 3 alpha. The antisense compound is useful for the treatment of a diseases associated with glycogen synthase kinase 3 alpha such as diabetes, a neurological disorder, a haematopoietic disorder, a hyperproliferative disorder or a developmental disorder. The antisense compounds may also be used prophylactically to prevent or delay infection, inflammation or tumour formation. The present sequence is a phosphorothicate antisense oligonucletide targetted to human glycogen synthase kinase 3 alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; full length cDNA; cDNA synthesis; oligo-capping; PCR primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human cDNA clone-specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 2 A; 8 C; 2 G; 8 T; 0
                                                                                                                                                                                                                      Example 18; Page 129; 1380pp + Sequence Listing; English
                                                                                                                                                                                                                                                                      830 Primers useful for synthesizing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-JUL-2000; 2000EP-00114089
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 05-SEP-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EP1130094-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAK95023;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAK95023
                                                                                                                                                                                                                                                                                                                                            Wakamatsu
                                                                                                                                                                                                                                                                                                                                                                                             (HELI-) HELIX RES INST.
                                                                                                                                                                                                                                                       genetic manipulation.
                                                                                                                                                                                                                                                                                                                                                            H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4871
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sapiens
                                 provided
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        l Similarity
17; Conserv
                                                                                                                                                                                                                                                                                                                                                            Nishikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CTCAGTTTCTTTCTCTGCAA 4890
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                             2000JP-00118774
2000JP-00183765
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry
                                                                                                                                                                                                                                                                                                                                          Sugiyama
                                   ä
                                                                                                                                                                                                                                                                                                                                                                                                                                                                99JP-00194486
                                   the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.3%;
                                                                                                                                                                                                                                                                                                                                          Isogai T,
Na T, Nagai
                                                                                                                                                                                                                                                                                                                                     -1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 primer,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                          Hayashi K,
K, Kojima
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO:
                                                                                                                                                                                                                                                                       full length cDNA clones and their use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U; 0 Other;
                                                                                                                                                                                                                                                                                                                                            တ
                                                                                                                                                                                                                                                                                                                                            Ishii S,
S, Otsuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                            Kawai Y;
T, Koga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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Query Match Best Local S Matches 17

Local Similarity hes 17; Conserv

Conservative

0.3%;

; Score 15.2; D ; Pred. No. 9.3e 0; Mismatches

.3e+02; DB 1; Other;

Length 20;

Indels

0

Gaps

0

Sequence

20

B₽;

4 P,

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C; 5

գ; 5

H.

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RESULT 854
AAK95024
ID AAK950
XX AAK950
XX Human
XX Human
XX Human
XX Homo &
XX Homo &
YX EP113(
XX EP113(
XX Homo &
YX Human
XX Human
XX Human
XX Human
YX OS-SEI
XX Human
YX Human
RESULT 855
AAC87238/c
ID AAC872
XX
AC AAC872
XX
AC AAC872
XX
DT 09-MAR
XX
DE Phosph
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                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates to primers for synthesising full length cDNA clones. 830 cDNA molecules encoding a human protein have been isolated and nucleotide sequences of 5'- and 3'-ends of the cDNA molecules have been determined. Primers for synthesising the full length cDNA are useful for clarifying the function of the protein encoded by the cDNA. The full length clones were obtained by construction of full length enriched cDNA libraries that were synthesised by the oligo-capping method. The primers enable the production of the full length cDNA easily without any special
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20 BP; 4 A; 6 C; 5 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JUL-1999; 99JP-00194486.
11-JAN-2000; 2000JP-00118774.
02-MAY-2000; 2000JP-00183765.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP1130094-A2
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                                                          09-MAR-2001
                                                                                                                                                                       AAC87238 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2001-524255/58.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-JUL-2000; 2000EP-00114089.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    05-SEP-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; full length cDNA; cDNA synthesis; oligo-capping; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAK95024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAK95024 standard;
  Phosphorothicate poly T oligonuclectide,
                                                                                                                  AAC87238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  830 Primers useful for synthesizing full length cDNA clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (HELI-) HELIX RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human cDNA clone-specific primer, SEQ ID NO: 4269
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3593 TTGCTCAGGCTAATCTCAAA 3612
                                                                                                                                                                                                                                                                                                                                                                       3593 TTGCTCAGGCTAATCTCAAA 3612
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1s. The present sequence is a primer used to amplify a human provided in the invention
                                                                                                                                                                                                                                                                                                                     Ľ
                                                                                                                                                                                                                                                                                                                                                                                                                                                         h 0.3%;
Similarity 85.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nishikawa T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18; Page 129; 1380pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                     TIGCCCAGGCTAGTCTCGAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TTGCCCAGGCTAGTCTCGAA 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             manipulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sugiyama
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Isogai T,
na T, Nagai
                                                                                                                                                                             20
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                                                                                                                                                                             ВP
                                                                                                                                                                                                                                                                                                                                                                                                                               0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.2; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hayashi K, Is
K, Kojima S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                               No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            9.3e+02;
     SEQ ID NO:17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ishii S,
S, Otsuki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kawai Y;
T, Koga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Koga H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and their use
                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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RESULT 856 AAC87230/c

AAC87230 standard; DNA; 20

ВP

AAC87230

문 8

20

Matches Query Match

5393 AAAAAAATACAAAAAAGAAA 5412

Local

Similarity

0.3%;

Score 15.2; DB 1; Pred. No. 9.3e+02; Mismatches

DB 1; Length 20; Indels

0

Gaps

0

0

Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;

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The invention relates to the use of an immunostimulatory single-stranded CC DNA-binding protein in screening assays to identify compounds which bind CC to it and thereby act as functional modifiers of immunostimulatory CC oligodeoxynucleotide (DDN) activity. Such modifiers of ODN activity CC consist of immunostimulatory DNA binding inhibitors, immunostimulatory CDNA mimics, and immunostimulatory DNA binding inhibitors, immunostimulatory CDNA mimics, and immunostimulatory DNA binding competitors, and to optimize an immunostimulatory DNA-binding proteins can also be used in screening CC complexes of an immunostimulatory DNA-binding protein bound to an CC immunostimulatory DNA can additionally be used to screen a panel of complexes of an immunostimulatory DNA-binding protein bound to an CC immunostimulatory DNA can additionally be used to screen a panel of complexes of an temptode of the invention are the RNA-binding proteins CC the immunostimulatory DNA. The immunostimulatory DNA-binding proteins CC uncleolin, bnRNP D, AUF1, bnRNP Al and lupus La protein. The screening CC mucleolin, bnRNP D, AUF1, bnRNP Al and lupus La protein. The screening CC methods are useful for identifying a compound that inhibits interaction CC between immunostimulatory DNA and an immunostimulatory DNA-binding protein and for identifying agonists useful in immunostimulatory DNA cellular target complex is useful in screening for immunostimulatory DNA cellular target complex investigation of structure/scriviy relationships of immunostimulatory CDNA cellular target complex investigation of structure/scriviy relationships of immunostimulatory CDNA cellular target complex in the complex in an oligonucleotide used in an exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Immunostimulatory DNA binding proteins to identify immunostimulatory DNA functional modifiers, immunostimulatory DNA binding competitors and to optimize immunostimulatory oligodeoxynucleotides for stimulation.
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03-MAR-2000; 2000US-0186845P
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09-MAR-2001

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Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA mimics, and immunostimulatory DNA agonists and antagonists.

Immunostimulatory DNA-binding proteins can also be used in screening methods to identify immunostimulatory DNA binding competitors, and to optimize an immunostimulatory ODN for immune stimulation. Isolated complexes of an immunostimulatory DNA-binding protein bound to an immunostimulatory ODN can additionally be used to screen a panel of candidate target molecules to identify the cellular target molecules of the immunostimulatory ODN. The immunostimulatory DNA-binding proteins used in the methods of the invention are the RNA-binding proteins nucleolin, hnrNP D, AUFI, hnrNP AI and lupus La protein. The screening methods are useful for identifying a compound that inhibits interaction between immunostimulatory DNA and an immunostimulatory DNA-binding protein sueful in screening for immunostimulatory DNA-binding protein and for identifying agonists useful in immunotherapy. The complex is useful in screening for immunostimulatory DNA cellular target
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA-binding protein in screening assays to identify compounds which bind to it and thereby act as functional modifiers of immunostimulatory oligodeoxynucleotide (DDN) activity. Such modifiers of DDN activity consist of immunostimulatory DNA binding inhibitors, immunostimulatory
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                                                                                                                                                                                                                                                                                                                                      molecules. The candidate immunostimulatory ODN competitors allow the investigation of structure/activity relationships of immunostimulatory DNA-binding proteins and immunostimulatory ODNs. The present sequence represents an oligonucleotide used in an exemplification of the inver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunostimulatory DNA binding proteins to identify immunostimulatory functional modifiers, immunostimulatory DNA binding competitors and toptimize immunostimulatory oligodeoxynucleotides for stimulation.
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03-MAR-2000; 2000US-0186845P.
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                                                                                                                                                                                                                                                                      Sequence 20 BP; 0 A; 0 C; 0 G;
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                                                                                                                                      Conservative
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Pred. No. 9
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                                                                                                                                   Indels
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RESULT 857 AAC87241/c ID AAC87241 standard; DNA; 20 BP

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Query Match Best Local S Matches 17

Local Similarity

0.3%;

Score 15.2; DB 1; Pred. No. 9.3e+02; Mismatches

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5393 АЛАЛАЛАТАСЛАЛАЛАСЛАЛ

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The invention relates to the use of an immunostimulatory single-stranded CDNA-binding protein in screening assays to identify compounds which bind CC to it and thereby act as functional modifiers of immunostimulatory CC oligodeoxynucleotide (ODN) activity. Such modifiers of of DNA activity CC consist of immunostimulatory DNA binding inhibitors, immunostimulatory CDNA mimics, and immunostimulatory DNA binding inhibitors, immunostimulatory CDNA binding proteins can also be used in screening CC methods to identify immunostimulatory DNA binding competitors, and to optimize an immunostimulatory DNA binding protein bound to an immunostimulatory DNA-binding protein bound to an immunostimulatory DNA-binding protein bound to an CC candidate target molecules to identify the cellular target molecules of CC candidate target molecules to identify the cellular target molecules of CC candidate target molecules are the knA-binding proteins CC used in the methods of the invention are the knA-binding proteins CC mucleolin, harny D, AUPI, harny Al and lupus La protein. The screening CC methods are useful for identifying a compound that inhibits interaction between immunostimulatory DNA and an immunostimulatory DNA-binding proteins and for identifying agonists useful in immunostimulatory DNA cellular target CC investigation of structure/activity relationships of immunostimulatory CDNA-binding proteins and immunostimulatory DNA cellular target CC investigation of structure/activity relationships of immunostimulatory CDNA-binding proteins and immunostimulatory DNA centerion of the invention CC the invention of structure/activity relationships of immunostimulatory CDNA-binding proteins and immunostimulatory DNA centerion of the invention CC the invention of structure/activity relationships of immunostimulatory CDNA-binding proteins and immunostimulatory DNA centerion of the invention CC immunostimulatory DNA centerion of the invention CC immunostimulatory DNA centerion of immunostimulatory DNA centerion of immunostimulatory DNA cente
Sequence 20
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                                                   represents an oligonucleotide used in an exemplification
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03-MAR-2000; 2000US-0186845P.
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cellular target identification; oligonucleotide optimisation;
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BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                        the invention
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RESULT 858
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequence represents a synthetic DNA template molecule used to a demonstrate the method of the invention. The invention relates to a method of modifying (e.g. 3' end labelling with 32P dATP) the 3' terminus of an RNA molecule by providing a DNA oligonucleotide, complimentary to the 3' end of the RNA molecule, with an overhang at the 5' end which allows incorporation of the labeling nucleotide into the RNA molecule. The method, based on the synthesis of okazaki fragments, is useful for labeling and modifying the 3'-termini of okazaki fragments, is useful for labeling and modifying the 3'-termini of other nucleic acids such as DNA fragments. The method is a simple and efficient way of labeling or modifying RNA 3'-termini using DNA polymerase and a synthetic template
Scaffold protein; antibody mimic; fibronectin type III domain; randomised loop; randomised beta-sheet; diagnostic purpose; protein designing; probe; tenth module of human Fn3; 10Fn3; fibronectin module of type III; Fn3; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Modifying a 3' terminus of a pre-selected DNA sequence, useful for labeling and modifying 3'-termini of other nucleic acids, comprises a synthetic nucleotide template with a defined overhang nucleotide
                                                                                                                                                                                                                                                                                                                              AAD16997 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (CHEN/) CHEN G.
(HUAN/) HUANG Z.
(SZOS/) SZOSTAK J W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17-OCT-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                  probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        overhang nucleotides
                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-00173936.
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                                                                                                                                                                                                                                                                                                                                 ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to an array of proteins (antibody mimics) comprising a fibronectin type III domain having a randomised loop, a randomised beta-sheet, or their combination, and has the capacity to bind to a compound that is not bound by a corresponding naturally-occurring fibronectin, immobilised onto a solid support. The antibody mimics is useful for detecting a compound preferably a protein, in a biological sample. It is also useful to detect one or more different analytes simultaneously in a sample. Hence is useful for diagnostic purposes. It is also useful for the purpose of designing proteins capable of binding to virtually any compound of interest. The present sequence is a capture probe used to self-assemble and anchor the tenth module of human fibronectin module of type III (Fn3) (10Fn3) which is used in an exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fibronectin scaffold protein array for obtaining a protein/compound which binds to a compound/protein, comprises a fibronectin type III domain having a randomized loop, a randomized beta-sheet or their combination.
                          28-JUL-1999;
                                                      28-ЛП-1999;
                                                                                                              DE19935302-A1
                                                                                                                                         Unidentified.
                                                                                                                                                                               Transport; membrane; cytostatic; virucide; vasotropic; dermatological; antipsoriatic; antiasthmatic; gene therapy; tumor cell; antisense;
                                                                                                                                                                                                                            Conjugate forming oligonucleotide ON5 SEQ
                                                                                                                                                                                                                                                        15-MAY-2001
                                                                                                                                                                                                                                                                                    AAF60896,
                                                                                                                                                                                                                                                                                                                AAF60896 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 41; 67pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-557782/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      29-FEB-2000; 2000US-00515260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-FEB-2001; 2001WO-US006414.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOTAHA (-TAHA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     $ . . . .
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                              5393 АЛАЛАЛАТАСЛАЛАЛАGAЛА 5412
                                                                                                                                                                   therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                          l Similarity
17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                        (first
                                                                                                                                                                 antiasthmatic; gene t
drug; phosphodiester
                         99DE-01035302
                                                        99DE-01035302
                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         85.0%;
                                                                                                                                                                                                                                                        entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.3%;
                                                                                                                                                                                                                                                                                                                20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               English.
                                                                                                                                                                                                                                                                                                                ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15.2;
Pred. No. 9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        9.3e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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(AVET) AVENTIS PHARMA DEUT GMBH.

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RESULT 861
AAH5699
ID AAH569
XX AAH569
AC AAH569
XX 10-SEE
XX 10-SEE
XX UIgani
KW Ligani
KW glucoo
KW MR; pe
KW progee
KW thyroj
KW transes
                                                                                                                                                                                                                                                                                                                                                                                                                                                    S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC Ar-(XY-R-I) n (II). Ar = group containing at least one argument of the containing at least one argument ring;

CX = O or N (Sic); Y = O, S or NH-R 2 (sic); R I = optionally substituted

CY = 3C alkyl (optionally containing double and/or triple bonds); R = 1

CY optionally substituted 1-18C alkyl (optionally containing double and/or triple bonds); n = integer of 1 or more. (A) is bonded to (B) directly or cy via a chemical group, provided that the chemical group is other than CH 2

CY = Sif the bond is via a phosphodiester linkage of (A). The invention also consiste the construction of the preparation of a conjugate (I') of (A') a molecule to be transported and (B') at least one aryl residue (not restricted to cy of (II)), by preparing (A') containing a reactive function at the position can which (B') is to be bonded, preparing (B') and reacting (A') and (B'); and (ii) the use of aryl groups (II) (optionally bonded via a chemical) cy open (II), by preparing (A) across biological membranes. The products of the invention have cytostatic, virucide, vasotropic, dermatological.

CY antipporiatic and antiasthmatic activity and can be used for gene therapy. Conjugation of (A) with (B) is useful for transporting (A) across biological membranes or into eukaryotic can be used for gene cytostatically bacterial, yeast or mammalian cells, including human cells, particularly tumor cells). Medicaments, diagnostic agents and test kits conjugated viral infections or disease associated with integrins or cells. (Cy optionally (I) are antisense or labeled oligonucleotides for in vivo diagnostic use, e.g. by in situ or cells interactions (e.g. restenosis, vitiligo, psoriasis or asthma); or hybridization. Conjugates for invivo diagnostic use, e.g. by in situ or conjugates (I) are finorescently labeled, allowing (I) is superior to that obtained using other conjugated groups related to (II); e.g. oligonucleotides conjugated with fluorescein diacetate (within the scope of (B)) have superior uptake to corresponding fluorescein
                                                                                                                                                                                                                                                                                                                                                                                                                    밁
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Best Local S
Matches 17
glucocorticoid receptor protein; GR; mineralocorticoid receptor protein; GR; mineralocorticoid receptor protein; GR; mineralocorticoid receptor protein; proxessore protein; proxessore progesterone receptor protein; PR; pregnane X receptor protein; PXR; thyroid hormone receptor protein; TR; vitamin D receptor protein; VI transactivation; ERalpha; breast cancer; PCR primer: Proham of the protein; VI
                                                                                                                                                                                                                  10-SEP-2001
                                                                                                                                                                                                                                                                                                       AAH56999 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         molecule to be transported and (B)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New substituted aryl conjugates of parent molecules, especially oligonucleotides, having improved transmembrane and intracellular transport properties, useful as medicaments or diagnostic agents.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Uhlmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                              5393 AAAAAAATACAAAAAAGAAA 5412
                                                                                                                                                                    oestrogen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                    receptor alpha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
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                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.2; DB 1; Length 20; Pred. No. 9.3e+02;
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                                                                                                                                                                      search PCR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         conjugate (I) which consists of (A) a B) at least one aryl residue of formula
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gothe
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                                                                                                                                                                    primer
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RESULT 862
AAS63428
ID AAS634
XX AAS634
XX AAS634
XX Oligor
XX Synthe
                                                                                                                                                                                                                                                                                                                                                                                                     CC peroxisome proliferator-activated receptor protein (MR),

CC receptor protein (PR), pregname X receptor protein (PRA), progesterone

CC receptor protein (PR), pregname X receptor protein (PXR), thyroid hormone

CC acids encoding them and cells comprising them and a specified reporter

CC gene for the ligand dependent transcriptional factor. These proteins are

CC useful in the modulation of ligand dependent transcriptional factor

CC activity. The cells, mutant ERalpha and the polymuclaotide encoding it

CC may be used in assays for qualitatively analysing an activity for

CC mutant ligand dependent transcriptional factors, for screening

CC mutant ligand dependent transcriptional factors, for revaluating an

CC activity for transactivation of a reporter gene by a test ERalpha and/or

CC for screening a compound useful for treating a disorder of a mutant

CC ERalpha, especially breast cancer
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                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
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21-JUL-2000; 2000JP-00220508.
02-AUG-2000; 2000JP-00234053.
03-AUG-2000; 2000JP-00235460.
03-AUG-2000; 2000JP-00235461.
03-AUG-2000; 2000JP-00235463.
                                                                                                                                                        AAS63428
                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ligand dependent transcriptional factors, nucleic acids encoding them and cells comprising them and a specified reporter gene, useful for screening agents for the treatment of breast cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present invention relates to ligand dependent transcriptional factors including oestrogen receptor (ER) alpha and beta protein, glucocorticoid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Saito K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             14-JUN-2001.
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                                                                                                                         29-JAN-2002
                                                                                                                                                                                      AAS63428 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 8; Page 214; 276pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                07-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-DEC-2000; 2000WO-JP008553
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                                                                                                                                                                                                                                                                                                                                                                                        BP; 6 A; 6 C; 3
                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                         (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99JP-00370667
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                                                                                                                                                                                      DNA;
                                                                                                                         entry)
                                                                                                                                                                                                                                                                                                                                          0.3%;
                                                                                                                                                                                      20
                                                                                                                                                                                      ВP
                                                                                                                                                                                                                                                                                                                                                                                        G; 5 T; 0
                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                          Score 15.2;
Pred. No. 9.
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                                                                                                                                                                                                                                                                                                                                        .3e+02
                                                                                                                                                                                                                                                                                                                                                                                        0 Other;
                                                                                                                                                                                                                                                                                                                                                           B
                                                                                                                                                                                                                                                                                                                                                          1; Length 20;
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Oligonucleotide-nanoparticle probe; diagnostic; nucleic acid detection; nanostructure; biochip;

forensic analysis;
biofilter; drug delivery;

Gaps

0

Oligonucleotide-nanoparticle probe #52.

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RESULT 863
AAF28481
ID AAF284
XX AAF284
XX AAF286
XX O3-AF8
DT 03-AF8
DX Random
XX Nuclei
KW diseas
KW cell l
XX
                                                                                                                                                                                                                                                                                                                                                                                   밁
                                                                                                                                                                                                                                                                                   S
                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to a method for detection of nucleic acid (I) having at least 2 portions, comprising treatment with nanoparticles that carry oligonucleotides complementary to at least 2 parts of (I), where detectable change caused by hybridisation of the oligonucleotide to (I) is observed. The method is used to detect (or to separate) specific (I), e.g. for diagnosing a wide variety of diseases, sequencing, in forensic analysis etc., and generally to detect analytes other than (I). The oligonucleotide-derivatised nanoparticles are also useful for preparing nanostructures useful, for example, as biochips, biofilters, mechanical devices, separation membranes, chemical sensors, in computers, and for drug delivery. Very stable nanoparticle-oligonucleotide conjugates can be produced, allowing their direct use (as probes) in polymerase chain reaction, i.e. they survive multiple heating/cooling cycles so do not need to be added after amplification. (I) are detected by simple colour change, without the need for special equipment, making possible rapid field testing for e.g. pathogens. AAS63374-AAS63448 represent
                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                      Matches
                               Nucleic acid detection; nanoparticle-oligonucleotide conjugate; disease diagnosis; forensic analysis; DNA sequencing; paternity testing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Detecting and separating nucleic acid, useful e.g. for diagnosis, comprises reaction with nanoparticles that carry oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-MAR-2000;
26-APR-2000;
                                                                             Random oligonucleotide,
                                                                                                              03-APR-2001
                                                                                                                                             AAF28481;
                                                                                                                                                                          AAF28481 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 18; Page 158; 404pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mirkin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-MAR-2001; 2001WO-US010071.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  complementary to parts of the target.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-656926/75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200173123-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NANO-) NANOSPHERE INC
                                                                                                                                                                                                                                                                                     5393 AAAAAATACAAAAAAGAAA 5412
                  line authentication;
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                                                                                                                                                                                                                                                                                                                                                                                                                  the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; 2000US-0213906P.
; 2000US-0254392P.
; 2000US-0255235P.
; 2001US-00760500.
; 2001US-00820279.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Letsinger RL,
Park S, Li Z;
                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000US-0200161P.
2000US-00603830.
                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                 0.3%;
                                                                              SEQ
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                                                                              IJ
                                                                                                                                                                                                                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                   Score 15.2; DB 1; Length Pred. No. 9.3e+02;
                therapy; ss
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                                                                              53
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                                                                                                                                                                                                                                                                                                                      Indels
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AAS10371

ID AAS

XX AAS

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DT 24--

XX Oli.

XX Nann

KW DNA

KW DNA

KW Eor

XX Syn

XX Key

FH Key

FT mis
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                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 864
                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is an oligonucleotide used in a method for detecting a nucleic acid having at least 2 portions. The method comprises hybridising the nucleic acid with oligonucleotides, such as the present sequence, attached to a substrate and/or particle and detecting a change in colour, conductivity or optical density. The method is useful for the diagnosis and/or monitoring of diseases, in forensics, in DNA sequencing, for paternity testing, for cell line authentication and for monitoring gene therapy. Detecting nucleic acids based upon observing a colour change is cheap, fast, simple, and does not require specialised or expensive equipment. The nanoparticle oligonucleotide conjugates remain stable for at least 6 months. A single base mismatch and as little as 20 femtomoles (fM) of target can be detected using the conjugates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Detecting nucleic acid, useful for e.g. diagnosis of diseases, for and DNA sequencing, comprises observing detectable change brought by hybridization of nucleic acid with substrate or particle bound
          misc_feature
                                                      Synthetic
                                                                               Nanoparticle; cyclic disulphide-oligonucleotide; DNA detection; DNA isolation; genetic disease; bacterial disease; viral disease; forensic science; paternity testing; gene therapy; ss.
                                                                                                                                          Oligonucleotide-cyclic disulphide linker,
                                                                                                                                                                      24-OCT-2001
                                                                                                                                                                                                   AAS10371
                                                                                                                                                                                                                                AAS10371 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 199; 205pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 2001-061976/07.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25-JUN-1999; 99US-00344667.
26-APR-2000; 2000US-0200161P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Taton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (MUCI/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MIRK/) MIRKIN C A.
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STORHOFF J J.
ELGHANIAN R.
TATON T A.
                                                                                                                                                                                                                                                                                                                                                            l Similarity
17; Conserv
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MUCIC R C.
                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Letsinger
                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                        (first
                           Location/Qualifiers
                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                          0.3%;
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                                                                                                                                                                                                                                ВP
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                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                           Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Storhoff JJ,
                                                                                                                                           α.
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                                                                                                                                                                                                                                                                                                                                                                                          20;
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RESULT 865
AAF99427/C
ID AAF994
XX AAF994
XX AAF994
XX AAF994
XX Immunc
XX Uaccin
XW Immunc
XW Immunc
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC may be coupled with colloidal gold particles (nanoparticles) and used to complete with colloidal gold particles (nanoparticles) and used to colloidal gold particles (nanoparticles) and used to colloidal gold) are used to both isolate and detect (e.g. by linking it to 2 or more complimentary nucleotides which have a nanoparticle attached to their 5' ends. The nanoparticles (e.g. colloidal gold) are used to both isolate and detect (e.g. by linking the comparticle to a fluorescent probe) the resultant complex. The methods are useful for detecting nucleic acids, natural or synthetic, and modified or commodified. The methods may also be applied in the diagnosis of genetic, bacterial and viral diseases, in forensics, in DNA sequencing, for cell line authentication, and for monitoring gene compartories in DNA sequencing, in the field to detect the presence of specific pathogens, for quick identification of an infection to assist in Cd drug prescription, and in homes and health centres for inexpensive first-curies exceening. The methods, which are based on observing colour change with the naked eye, are cheap, fast, simple, robust (reagents are considered are specialised or expensive equipment, and little or considered and an alternative for inexpensive first stable), do not require specialised or expensive equipment, and little or
                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 17
Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting a nucleic acid useful in e.g. diagnosing genetic, bacterial or viral diseases, by contacting the nucleic acid with oligonucleotides attached to nanoparticles and having sequences complementary a portion of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13-JAN-2000; 2000US-0176409P.
26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-00603830.
12-JAN-2001; 2001US-00760500.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 24; Fig 44;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           viral diseases, by contacting attached to nanoparticles and the nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-JAN-2001; 2001WO-US001190
                                                                                                                                                       AAF99427;
                                                                                                                                                                                          AAF99427 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The sequence represents a cyclic disulphide linked oligonucleotide which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200151665-A2
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                                                                         Immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         instrumentation is required
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CA,
                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                    AAAAAAAAAAAAAAAAAA 20
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Li Z;
                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "A is covalently linked to a cyclic-disulphide
                                                                                                                                                                                          DNA;
                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      323pp; English.
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                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                             Score 15.2;
Pred. No. 9
                                                                             #543
                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Storhoff
                                                                                                                                                                                                                                                                                                                                          9.3e+02;
hes 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                              Gaps
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RESULT 866
AAF9909/c
ID AAF990/c
XX AAF990
XX INMUND
XX INMUND
DI ILIMUND
XX VACCIN
KW VACCIN
KW INMUND
KW INTELL
KW INTELL
KW INTELL
XX SYNTHE
XX WO2001
XX SPAPR
PD 05-APR
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Best Local Simi
Matches 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a method for stimulating an immune response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present sequence is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothioate backbone
                                                                                                                Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothioate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vaccinating against tumors, infectious using immunostimulatory Py-rich and TG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-SEP-1999; 99US-0156113P.
27-SEP-1999; 99US-0156135P.
23-AUG-2000; 2000US-0227436P.
                                                  WO200122972-A2
                                                                                                                                                                                                 Immunostimulatory nucleic acid #215
                                                                                                                                                                                                                                  12-JUN-2001
                                                                                                                                                                                                                                                                                                    AAF99099
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Krieg AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (IOWA ) UNIV IOWA RES FOUND. (COLE-) COLEY PHARM GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200122972-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infectious disease; allergy; immune deficiency; phosphorothioate;
                                                                                                                                                                                                                                                                                                                                                                                                            5393 AAAAAAATACAAAAAAGAAA 5412
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                                                                                                                                                                                                                                                                                                                                                                                     20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schetter C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000WO-US026383.
                                                                                                                                                                                                                                  (first
                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.3%;
                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.2; DB 1
Pred. No. 9.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diseases, allergies and asthma nucleic acids.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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0,

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RESULT 867
AAF99431
ID AAF994
XX AAF994
XX Immuno
XX Vaccin
KW Vaccin
KW immuno
KW immuno
KW infect
XX Synthe
XX 100-APF
XX 25-SEI
PR 23-SEI
PR 23-AUI
PA (IOWA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        片
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Best Local S
Matches 17
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27-SEP-1999;
23-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       response. The method comprises administering an immunostimulatory nucleic acid to a non-rodent subject in sufficient quantity to stimulate an immune response. The present set is one such immunostimulatory nucleic acid. The immunostimulatory nucleic acid. The immunostimulatory nucleic acid. The immunostimulatory nucleic acids can be pyrimidine rich (py-rich) or thymidine (T) rich. The method is used to vaccinate subjects against tumour antigens, viral antigens (e.g. herpesviridae, retroviridae and/or orthomyxoviridae), bacterial antigens (e.g. toxoplasma, haemophilus, campylobacter, clostridium, Escherichia coli and/or staphylococcus), fungal antigens and/or parasitic antigens. The method is also useful for preventing cancer, asthma, infectious disease, allergy or immune deficiency. The present sequence can also be used to redirect a Th2 to a Th1 immune response and to activate immune cells. Note: the present sequence may have a phosphorothioate backbone
                                                                                                                                                                                                                                                         Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection; fungal infection; parasitic infection; cancer; asthma; infectious disease; allergy; immune deficiency; phosphorothicate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vaccinating against tumors, infectious diseases, allergies and using immunostimulatory Py-rich and TG nucleic acids.
                                                                     25-SEP-1999;
27-SEP-1999;
                                                                                                                       25-SEP-2000; 2000WO-US026383.
                                                                                                                                                         05-APR-2001.
                                                                                                                                                                                           WO200122972-A2
                                                                                                                                                                                                                                                                                                                                             Immunostimulatory nucleic acid #547.
                                                                                                                                                                                                                                                                                                                                                                                12-JUN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                  AAF99431;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAF99431 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 101; Page 42; 338pp; English.
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                                                       23-AUG-2000;
   (IOWA ) UNIV IOWA RES FOUND. (COLE-) COLEY PHARM GMBH.
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                                                   99US-0156113P.
99US-0156135P.
2000US-0227436P.
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99US-0156135P.
2000US-0227436P.
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RESULT 868
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AC AAF995
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Best Local &
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                                                                                                                                                                                                                                                                                                                      25-SEP-1999; 99US-0156113P
27-SEP-1999; 99US-0156135P
23-AUG-2000; 2000US-0227436P
                                                                                                                   WPI; 2001-273485/28.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             fungal infection; parasitic infection; cancer; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vaccine; cytostatic; virucidal; bactericidal; fungicidal; anti-parasitic; immunostimulatory; tumour; viral infection; bacterial infection;
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Pred. No. 9.3e+02;
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Vaccinating against tumors, infectious diseases, allergies using immunostimulatory Py-rich and TG nucleic acids.

and asthma

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RESULT 869
AN437989
ID AN437989
XX AN4379
XX SNP 8p
XX SIngle
KW Single
KW Single
KW Lesch-
KW GOLLesch-
KW Inflam
XX Homo 8
XX HOMO 9
XX HOMO 8
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Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Single nucleotide polymorphism; SNP; single nucleotide primer extension; SNPE; genotyping; agammaglobulinaemia; diabetes insipidus; cancer; Lesch-Nyhan syndrome; muscular dystrophy; familial hypercholesterolaemia; polycystic kidney disease; osteogenesis imperfecta; autoimmune disease; acute intermittent porphyria; rheumatoid arthritis; multiple sclerosis; inflammation; forensic investigation; paternity analysis; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                             New genotyping oligonucleotide, useful for detecting the presence, absence or identity of single polynucleotide polymorphism in a nuc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Picoult-Newburg L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP specific upper PCR primer SEQ ID 785.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-OCT-1999;
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Pred. No. 9.3e+02;
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Sequences AAH37205 - AAH40944 represent PCR primers, single nucleotide primer extension (SNPE) primers, and the sequences of regions flanking sites of single nucleotide polymorphisms SNPs. The present invention includes kits for determining the presence or absence of a SNP, using the oligonucleotides of the invention. The PCR primers are used to amplify a

sequence,

SNPE primer is used as a genotyping

primer.

This invention describes a novel DNA sequence (I) encoding the MTR1 protein that: (i) has at least one biological activity of a TRP (transient receptor potential) family protein; (ii) is connected with etiology of BMS (Beckwith-Wiedemann syndrome) and/or (iii) is connect.

18 connected

Page 19;

46pp;

German

sample

54; 83pp; English

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RESULT 870
AAH20526/c
ID AAH2055
XX AAH205
XX AAH205
XX Human
XX HIP15
XW Lransi
XW Lransi
XW INTR1;
XW INTR1;
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XX HOMO (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Loca
Matches
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MTR1; TRP-related protein; Ca2+ regulation; calcium regulation; tumor; transient receptor potential family; BWS; Beckwith-Wiedemann syndrome; 11p15.5 abnormality; chromosome 11; anchorer; developmental activity; intracellular calcium ion regulation; hormone; growth factor; apoptosis; cell growth; cell death; cell differentiation; urogenital disease; polycystic kidney disease; calcium influx; Wilms tumor; rhabdoid tumor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               diseases, including, rheumatoid arthritis, mutury occavior, inflammation, cancer, nervous system diseases and infection by pathogenic microorganism. The method is also useful in forensic investigations and microorganism.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human
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                                                                                                                                                                                                                                                                                                    DNA encoding MTR1 protein, useful syndrome and tumors, also related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prawitt D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-NOV-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYGU-) UNIV GUTENBERG JOHANNES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pelletier J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PCR primer; ss
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Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B
                                                                                                                                                                                                                                                                                                    e.g. for treating Beckwith-Wiedemann proteins and antibodies.
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CC invention have anticancer and developmental activity. MRT1 is involved in CC regulation of intracellular calcium ion levels, which are essential for CC cellular responses to hormones and or growth factors; also in apoptosis CC and cell growth, death and differentiation, and in urogenital diseases, CC including polycystic kidney disease. (I) and related ribozymes, antisense CC RNA, proteins and antibodies (Ab)) are used to treat or prevent diseases CC associated with altered expression of the MRT1 gene or activity of its CC protein, or with calcium influx into cells, e.g. BWS, Wilms tumor, CC thabdoid tumors and rhabdomyosarcoma. Probes from (I), or Ab, are also composed for diagnosis of such diseases. (I) can also be used for recombinant CC production of MRT1 proteins (II) (used for analysis, characterization and therapy), as tissue or chromosomal markers, for identifying genetic CC diseases and related sequences, as primers for genetic fingerprinting, as source of oligonucleotides for biochips, and to raise anti-protein or CC compotitive assays for (II) are used to raise Ab, as reagents in CC compotitive assays for (II), as tissue markers; for identifying contents and in screening for (anti-approximation) and the method of the invention of the human MTR1 gene CC described in the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 17; Conserv
                    Preparing sulfurized 2' substituted phosphorothicate oligonuclectides useful in biological research, comprises phosphitylating the 5'-hydro: of a nucleic acid having a nucleoside with a 2' modification.
                                                                                                                                                                                                                                                           11-JAN-2000; 2000US-00481486
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Phosphorothioate; anti-viral therapy; stereochemical pathway;
                                                                                                                                                                                 (ISIS-)
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                                                                                                                                                                                 ISIS PHARM INC
                                                                                                                                         Ravikumar VT,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            'note= "All bases are phosphorothioate"
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                                                                                                                                         Cheruvallath
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Pred. No. 9.3e+02;
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                       The specification describes a method for identifying a nucleic acid in a pool of interest. The method comprises pooling individually identifiable nucleic acids into at least two pools of one nucleic acid each; expressing nucleic acid pools to obtain protein expression product pools; assaying protein expression product pools for products having interaction with target molecule; selecting nucleic acid pools corresponding to identified protein expression product pools; and identifying individual nucleic acids in identified nucleic acid pools. The method is useful for identifying a nucleic acid (e.g. cDNA) in a pool of interest and for functionally screening several nucleic acids. The method is also useful for screening genomic DNA libraries or other source of individual cDNAs, manners, synthetic libraries of nucleic acids e.g. combinatorial libraries. The present sequence was used in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a method for preparing phosphorothioate oligonucleotides having at least one nucleoside with a 2 modification. The method comprises phosphitylating the 5'-hydroxyl of a nucleic acid group having at least one nucleoside with a 2 modification in an acetonitrile. The present sequence was used to illustrate the method of the present invention. The method is useful for synthesising sulphurised 2' substituted phosphorothioate oligonucleotides, which may be used in molecular biological research, in applications such as anti-viral therapy, and for determining the stereochemical pathways of certain enzymes which recognise nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Screening nucleic acids (NA) in pool of interest comprises pooling, expressing NA to form expression product pool and identifying NA in NA pool corresponding to expression product pool having interaction with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Col 22; 19pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                7pp;
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Pred. No. 9.3e+02;
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RESULT 873
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AC AAP2
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                                                                                                                                                                CC S-Methythioura (DNmt) compounds to form triplexes with DNA oligomers. An CC increase in the C content of the oligos resulted in a large decrease in CC binding. This experiment was performed as an example of a method for CC preparing oligonuclectides comprising a backbone of alkyl or alkoxy CC thiourea linkages. The method is useful for preparing oligonuclectides CC for use in antisense or antisense therapy, to inhibit production of CC proteins associated with genetic diseases, cardiovascular, inflammatory CC and neurocellular diseases, and for antiviral therapy, e.g. to treat CC numan immunodeficiency virus, human-cytomegalovirus, influenza and herpes CC infections. The compounds are also useful as diagnostic reagents to CC detect the presence or absence of the target DNA or RNA sequences to CC which they specifically bind and by antagonising the normal biological CC cissue e.g. tissue differentiation, both in vivo and in ex vivo tissue CC cultures. The method provides an efficient and rapid solid-phase method CC cultures. The method provides an efficient and rapid solid-phase method CC for the synthesis of thiourea and S-methylthiourea
        Query Match
Best Local S
Matches 17
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11-DEC-1998;
02-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Preparing novel deoxymucleic alkyl thiourea oligonucleotide for use in antisense therapy, by synthesizing oligonucleotides comprising backbone of alkyl or alkoxy thiourea linkages in solution or on solid phase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Deoxynucleic g-Methythiourea; DMmt; antisense therapy; cardiovascular disease; inflammatory disease; neurocellular disease; antiviral therapy; human immunodeficiency virus; human-cytomegalovirus; influenza; herpes; infection; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence was used to demonstrate the ability of S-Methythiourea (DNmt) compounds to form triplexes with DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 7; Fig 16; 48pp; English.
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                                                                                                                             BP;
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98US-0111800P.
99US-00347443.
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                                                                                                                          A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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        Score 15.2; DB 1
Pred. No. 9.3e+02
0; Mismatches
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Pred. No. 9
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                                                              DB 1; Length 20;
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           Indels
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RESULT 874
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RESULT 875
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence is that of the antisense primer used in the PCR amplification of a 150 bp fragment of the human neurokinin NK-2 receptor gene. The sense primer is given in ANHS0608. Quantitative PCR was performed on total RNA extracted from breast cancer cells. The results showed that NK-2 expression is up-regulated in breast cancers. Agonists and antagonists that luence expression of NK-2 can be used for the treatment of breast cancer, bone marrow metastasis, pain, asthma, arthitis, aggressive behaviour and depression associated with NK-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NK-2; neurokinin receptor; PPT-I; human; preprotachykinin; cytostatic; analgesic; antiarthritic; antiasthmatic; antidepressant; breast cancer; metastasis; pain; arthritis; aggression; depression; haematopoietic disorder; gene therapy; PCR primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20
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                                                                                                                                                                                AAF23343 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BP; 4 A; 10 C; 0 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Oligonucleotide for

detection

of Mycobacterium

19-MAR-2001

(first entry)

AAF23343;

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RESULT 876
ABK47733
ID ABK477
XX
AC ABK477
XX
DT 18-JUN
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                                                                                                                                                                                                                                                         유
                                                                                                                                                                                                                                                                               The present sequence is an oligonucleotide developed using a Mycobacterium ITS (internal transcribed spacer region) nucleotide sequences. ITS DNA sequences from M. fortuitum, M. chelonae, M. abscessus, M. vaccae, M. flavescens, M. asiaticum, M. porcinum, M. acapulcensis, M. diernhoferi genes were identified. The oligonucleotides derived from these sequences were used to develop PCR primers and hybridisation probes for detection and identification of Mycobacterium. ITS has a more polymorphic region than 16S rRNA and also has a conserved region. It is polymorphic region than 16S rRNA and also has a conserved region. It is possible to develop pcr distinction of genotype. The oligonucleotide probes, attached to solid substrate, hybridise only with nucleotide sequences in ITS of specific mycobacteria, and thus they can detect and identify the specific mycobacteria by PCR amplification. Using the oligonucleotide primers or probes made of the specific topic bacteria, it is possible to detect mycobacteria, distinguish tuberculosis (TB) complex from non-tuberculosis mycobacteria
                                                                                                                                                                                              Query Match
Best Local S
Matches 17
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29-MAY-1999;
29-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel oligonucleotide sequences of internal transcribing spacer region non-tuberculosis mycobacteria (NTM) used as probes or primers for detecting and identifying mycobacteria and distinguish TB complex from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ITS; internal transcribed spacer region; Mycobacterium fortuitum; Mycobacterium chelonae; Mycobacterium abscessus; Mycobacterium vacc Mycobacterium flavescense; Mycobacterium assisticum; tuberculosis; Mycobacterium assisticum; tuberculosis; Mycobacterium acapulcensis; identification; Mycobacterium acapulcensis; identification; Mycobacterium diernhoferi; PCR primer; probe; detection; ss.
 18-JUN-2002
                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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                               ABK47733;
                                                           ABK47733 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAY-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-MAY-2000; 2000WO-KR000477
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (PARK/)
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                                                                                                                                                                1036 GAGTCACCCAACGCCCCAC 1055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2001-061527/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              36; Page 82; 89pp; English
                                                                                                                                     20
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KIM C M.
PARK H K.
                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                         20
                                                                                                                                     GAGTCACCGAACTCCCACAC 1
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                                                                                                                                                                                                                                                         BP;
                                                                                                                                                                                               Conservative
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 (first entry)
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99KR-00019632.
99KR-00019633.
99KR-00019634.
99KR-00019635.
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                                                                                                                                                                                                                                                         G; 6 T;
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Pred. No. 9.
                                                                                                                                                                                               Mismatches
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                                                                                                                                                                                                             9.3e+02;
                                                                                                                                                                                                                                                         0 Other;
                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                            Length
                                                                                                                                                                                               Indels
                                                                                                                                                                                              0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vaccae;
                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of.
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Beta-actin
                                                    PCR primer #3
                                                    used in invention relating
                                                     to TMOs
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Primer-dependent polymerase-mediated DNA synthesis; TMO; template-mimic oligonucleotide; nucleic acid amplification; multiplex RT-PCR; reverse transcriptase-PCR; Competimer method; PCR; primer; beta-actin; ss.

Unidentified

WO200218616-A1

30-AUG-2001; 2001WO-US027287.

01-SEP-2000; 2000US-0230184P.

HITB (HITB HITACHI CHEM CO LTD. HITACHI CHEM RES CENT INC.

Ke S;

WPI; 2002-315546/35.

oligonucleotide. dependent pefficiency Modulating g amplification efficiency of a target sequence in primer-polymerase-mediated DNA synthesis, useful for adjusting the y nucleic acid amplification comprises adding a template mimic-

Example; Page 11; 39pp; English.

oligonucleotide (TWO) to a primer-dependent polymerase-mediated DNA synthesis reaction mixture containing primers, to block a primer for amplifying a target sequence in the mixture from hybridising to the target sequence. The method is useful for adjusting the efficiency of target template nucleic acid amplification by controlling the ratio of template-like oligonucleotides to amplification by remember. The new method provides a convenient and efficient method for simultaneous detection of high and low-abundant genes in multiplex RT-PCR, and is more potent and easier to control than the Competimer method. The present sequence represents a PCR primer used in the example of the present invention The present invention relates to a method of modulating amplification efficiency of a target sequence in primer-dependent polymerase-mediated DNA synthesis. The method comprises adding a template-mimic

Sequence 20 BP; 5 A 9 C; 3 G; 3 T; 0 U; 0 Other;

Query Match Best Local & Matches Similarity Conservative 0.3%; 0 Score 15.2; UB 1, No. 9.3e+02 DB 1; Length 20; Indels 0 Gaps 0

吊 Ş 1262 GCCTACAGCCCCACCACCAC 1281 щ GGCTACAGCTTCACCACCAC

ABK97573

standard;

DNA;

20

₽P

07-OCT-2002 ABK97573; (first entry)

Human LCAT gene reverse PCR primer #20.

RESULT 877
ABK9753/c
ID ABK975
XX ABK975
XX O7-OCT
XX
DB Human
XX Lecith
XW Lecith
XW fish-e
KW popula
XW popula
XW popula
XW DS Homo 8 polymorphism; Lecithin-cholesterol acyltransferase; LCAT; Norum disease; gene therapy; fish-eye disease; atherosclerotic cardiovascular disease; forensic; population diversity; anthropological lineage; paternity testing; human; primer;

Homo sapiens

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RESULT 878
ABN84468/c
ID ABN844
XX ABN844
XX ABN844
XX CANDOX
DT 21-OCT
XX Carbox
XX C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present invention relates to a new polynucleotide comprising a nuclectide sequence which is a polymorphic variant of a reference sequence for lecithin-cholesterol acyltransferase (LCAT). The invention is useful for identifying an association between a trait (preferably a clinical response to drug targeting LCAT) and at least one genotype or chaplotype of LCAT gene. The method of the invention has applicability in developing diagnostic tests and therapeutic treatments for Norum disease, fish-eye disease and atherosclerotic cardiovascular disease. The chaplotyping and genotyping methods are useful for studying population diversity, anthropological lineage, the significance of diversity and lineage at the phenotypic level, paternity testing, forensic applications and for identifying association between the LCAT genetic variation and trait such as level of drug response or susceptibility to disease. In addition, the methods for identifying the LCAT haplotypes present in individuals are useful in the development of drugs targeting LCAT. For example, determining the frequency of individual LCAT haplotypes in a population with a specific disease, e.g. Norum disease, will facilitate the development of drugs targeting the LCAT isoform(s) that are most frequent in that disease population. The present nucleic acid sequence represents one of a collection (ABK97534-ABK97573) of PCR primers that the methods of the invention to detect polymorphisms in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local
                                                                                                                                                          Carboxypeptidase; islet of Langerhans; endocrine; stem cell; cell differentiation; diabetes; antidiabetic; cell therapy; enzyme; PCR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel isolated polymorphic variant polynucleotide of lecithin-cholesterol acyltransferase gene, useful for studying expression and biological function of the gene, and for therapeutic, diagnostic or forensic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chew A,
   WO200259278-A2
                                                                                                                                   primer;
                                                                                                                                                                                                                                                            Carboxypeptidase A forward PCR primer.
                                                                                                                                                                                                                                                                                                                               21-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                               ABN84468
                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABN84468 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 20 BP; 4 A; 9 C; 3 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               754 GGGCTGAGGTCACTGTGCGC 773
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                                                                                                                                   88.
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                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            20
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Pred. No. 9
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RESULT 879
ADD46625;
ID ADD46625;
XX ADD46625;
AC ADD4662
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    culture of undifferentiated embryonic stem cells, selecting endocrine culture of undifferentiated embryonic stem cells, selecting endocrine comparison cells (BPCs), expanding the BPCs by culturing in an expansion medium that comprises a growth factor, and differentiating the expanded BPCs in a differentiation medium. The present sequence is a forward PCR primer for carboxypeptidase A. Use with the primer given in ABN84469 comprises a product of 521 bp. A primer set (see ABN84464-71) was used in an example of the invention for RT-PCR analysis of endocrine pancreatic gene expression in mouse embryos at stages 1 and 5. The results suggested that the culture conditions used to generate pancreatic endocrine cells supported the differentiation of pancreatic cells. Pancreatic endocrine cells cells generated in vitro can be used as a model system to study the cells of insulin release, to study agents that affect insulin secretion, and for the development of artificial islet and artificial pancreas tissues for use in the treatment of pancreatic endocrine disorders, particularly type
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Differentiating embryonic stem cells from endocrine cells, for treating a pancreatic endocrine disorder, e.g. type I or type II diabetes, comprises culturing endocrine cells in an expansion medium containing a growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24-JAN-2001; 2001US-0264107P.
06-FEB-2001; 2001US-0266917P.
18-OCT-2001; 2001US-0344548P.
                                                                                                                                                                                                                                                                                                                                                                        Human ABCC11 intron20/exon21 junction site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 20
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                                                                                                                                                                                                                    Homo sapiens.
                                                                                                                                                                                                                                                                                cholesterol transport;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention
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                                                                                                                                                                                                                                                                                                         protein; paroxysmal kinesigenic choreoathetosis; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GGCATCAAACACACATTTGC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP; 4 A; 3 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Blondel O,
                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
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                                                                                     /*tag=
                            note= "partial"
                                                                                                                                                    Location/Qualifiers
                                                               number=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA; 20 BP
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                                                                                                                                                                                                                                                                             gene therapy; human; ds
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Pred. No. 9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9.3e+02
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one

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ARSULT 880
ABS77742/c
ID ABS7777
XX
AC ABS7777
XX
AC ABS7777
XX
DT 13-DEC
XX
Angiog
XX
Angiog
XW
Angiog
XW
Cornea
XW
Cornea
XW
Tuboux
XW
XX
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to novel ABCC11 nucleic acids and proteins. ABCC11 sequences are used in the manufacture of a medicament for treating and/or preventing subjects affected by paroxysmal kinesigenic choreoathetosis. They may be used for treating or preventing subjects affected by a dysfunction of the transport of anionic drugs such as GSH, glucuronate, or sulphate conjugated to acidic ligands such as GSH, glucuronate, or sulphate conjugated drugs. Compositions comprising the ABCC11 polypeptide may also be used in the treatment and/or prevention of a deficiency in the transport of cholesterol or inflammatory ligid substances and diseases mapped on the chromosome locus 16q12. ABCC11 protein can be used to treat pathologies linked to the transport of lipophilic substances. The invention is used in gene therapy. The present sequence is human ABCC11 intron/exon junction site
                                                                            tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma, wound granulation; intestinal adhesion; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 43; 118pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New ABCC11 nucleic acids and proteins, useful in manufacturing a medicament for treating and/or preventing paroxysmal kinesigenic choreoathetosis, or pathologies linked to the transport of lipophilic
                                                                                                                                                                                                                                                                                                                                                                                  ABS77742 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 5 A; 4 C; 6 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              choreoathetosis, or pathologies linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rosier-Montus M, Prad
Allikmets R, Denefle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-MAR-2001; 2001US-0272757P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-MAR-2002; 2002WO-EP003241
                        Synthetic
                                                                                                                                                                                                                Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth;
                                                                                                                                                                                                                                                       Angiogenesis inhibitory oligonucleotide #226.
                                                                                                                                                                                                                                                                                                   13-DEC-2002
                                                                                                                                                                                                                                                                                                                                          ABS77742;
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(USSH ) US
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US DEPT HEALTH & HUMAN SERVICES.
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/number= 2
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                                                                                                                                                                                                                                                                                                 entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 15.2; DB 1; Length Pred. No. 9.3e+02;
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RESULT 881
ABS78072/c
ID ABS780
XX ABS780
XX ABS780
XX ABS780
XX Angiog
XX Elemont
XX Cornea
XW Cornea
XW rubes
XW plaque
XX RW plaque
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX Synthe
XX I1-UUL
XX 14-DEC
XX 14-DEC
XX 14-DEC
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                      Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth; tumour metastasis; precancerous lesion; theumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to inhibiting angiogenesis in a subject, coadministering at least one antiangiogenic nucleic acid molecule included is a kit comprising a first container housing the antian
   14-DEC-2000; 2000US-025534P
                                     14-DEC-2001; 2001WO-US048458
                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                  Angiogenesis inhibitory oligonucleotide #556.
                                                                                                                                                                                                                                                                                                                                                                                                       13-DEC-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                             ABS78072;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABS78072 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Inhibiting angiogenesis in a subject, involves administering antiangiogenic nucleic acid molecule to the subject.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-566690/60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Page 23; 276pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAAAAAAAAAAAAAAAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention
                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          85.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.2; DB 1;
Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            at least
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            comprising
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RESULT 882
ABS78076
ID ABS780
XX ABS780
XX APSiog
XX Angiog
XX Angiog
XW Lumouz
XW Cornea
XW MO2002
XW MO2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    δ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma a hypertrophic scars. The present sequence is an antiangiogenic nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          administering at least one antianglogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antianglogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-DEC-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Inhibiting angiogenesis in a subject, involves administering at least one antiangiogenic nucleic acid molecule to the subject.
                                  WPI; 2002-566690/60
                                                                                           Bratzler RL;
                                                                                                                                                                                                                                                                                                                                                                                                    WO200253141-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  angiofibroma; wound granulation; intestinal adhesion; atherosclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth;
                                                                                                                                                                                                                                                                            14-DEC-2001; 2001WO-US048458.
                                                                                                                                                                                                                                                                                                                                           11-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Angiogenesis inhibitory oligonucleotide #560.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABS78076 standard; DNA; 20
                                                                                                                                                          (COLE-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2002-566690/60
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                                                                                                                                                       COLEY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAAAAAATACAAAAAAGAAA 5412
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             invention
                                                                                                                                                                                                                   2000US-0255534P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         hypertrophic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first
                                                                                                                                                          PHARM
                                                                                                                                                          GROUP INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         scar.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.2;
Pred. No. 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 T; 0 U; 0 Other;
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.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a subject, comprising cid molecule. Also ing the antiangiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20
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Inhibiting angiogenesis in a subject, involves admin antiangiogenic nucleic acid molecule to the subject.
                                                                                                               involves administering
                                                                                                               at least one
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Claim 2; Page 29; 276pp; English

administering at least one antianglogenic nucleic acid molecule. Also included is a kit comprising a first container housing the antianglogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumacoid arthritis, psociasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia, reposeds rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma and hypertrophic scars. The present sequence is an antiangiogenic nucleic The invention of the invention relates to inhibiting angiogenesis in a subject, comprising

Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

밁 S Best Local Similarity Matches 17; Conserv Query Match Best Local 5393 АЛАЛАЛАТАСАЛАЛАДАЛА 5412 AAAAAAAAAAAAAAAAAAA 20 Conservative 0.3%; <u>.</u> Score 15.2; Pred. No. 9. Mismatches 9.3e+02 멂 Length Indels 20; 0, Gaps <u>°</u>

ABS78288 Angiogenesis inhibitory oligonucleotide #772 13-DEC-2002 ABS78288, standard; (first entry) DNA; 20 ΒP

ABS78288/c
ID ABS7828
XX ABS7828
XX ABS782
XX ADS109
XX Angiog
XX Angiog
XX Angiog
XX Angiog
XX Angiog
XX Angiog
XX Plaque
XX Solero
XX Solero
XX Solero
XX Solero
XX I1-JUL
XX I1-JUL
XX I1-JUL
XX I1-JUL
XX I1-JUL
XX I1-JUL
XX III-JUL
XX III-J tumour metastasis; precancerous lesion; rheumatoid arthritis; psoriasis; diabetic retinopathy; retinopathy of prematurity; macular degeneration; corneal graft rejection; neovascular glaucoma; retrolental fibroplasia; rubeosis; Osler-Webber Syndrome; myocardial angiogenesis; plaque neovascularisation; telangiectasia; haemophiliac joint; angiofibroma; wound granulation; intestinal adhesion; atherosclerosis; scleroderma; hypertrophic scar. Angiogenesis inhibitor; ss; angiogenesis; solid tumour growth;

Synthetic

WO200253141-A2

11-JUL-2002

14-DEC-2001; 2001WO-US048458.

14-DEC-2000; 2000US-025534P.

Bratzler RL

(COLE-) COLEY

PHARM GROUP INC

2002-566690/60

antiangiogenic Inhibiting angiogenesis in a subject, antiangiogenic nucleic acid molecule (nucleic acid ö involves administering at least the subject. one

Claim 2; Page 33; 276pp; English

The invention relates to inhibiting nhibiting angiogenesis antiangiogenic nucleic in a s subject, c i molecule. comprising e. Also

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RESULT 884
ABL39402/c
ID ABL394
XX ABL394
XX ABL394
XX ABL394
XX ARTibo
KW ARTibo
KW ARTibo
KW ARTibo
KW ARTIF
PT RODIF
XX SYNTHE
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                                                                                                                                                                                                                                                                                                         Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   included is a kit comprising a first container housing the antiangiogenic nucleic acids, and instructions for administering them to a subject having a condition characterised by unwanted angiogenesis. The method is useful for inhibiting angiogenesis associated with solid tumour growth, tumour metastasis, precancerous lesion, rheumatoid arthritis, psoriasis, diabetic retinopathy, retinopathy of prematurity, macular degeneration, corneal graft rejection, neovascular glaucoma, retrolental fibroplasia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rubeosis, Osler-Webber Syndrome, myocardial angiogenesis, plaque neovascularisation, telangiectasia, haemophiliac joints, angiofibroma, wound granulation, intestinal adhesions, atherosclerosis, scleroderma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-JUN-2000; 2000US-0213346P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             angiogenesis; metastasis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     modified_base
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                                                                                                                                                                                                                                                                                        cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hartmann
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Pred. No. 9.3e+02
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The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for

Disclosure; Page 309; 312pp; English

treating or preventing cancer such as basal cell carcinoma, cancer, bone cancer, brain and central nervous system (CNS)

cancer,

cervical cancer,

cancer,

Query Match

Sequence

20

BP; 20 A; 0 C; 0 G; 0 T; 0 U;

0 Other;

Score 15.2;

DB 1; Length 20;

bladder

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RESULT 885
ABL38648
ID ABL386
XX ABL386
XX Immuno
XX Antibo
KW Antibo
KW angiog
XX Synthe
PN WO2001
XX 22-JUN
XX 22-JUN
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                                                                              The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, cesophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovarian cancer, pancreatic cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          tissue cancer, oesophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, non-Hodgkin's lymphoma, melanoma, myelancar cavity cancer, osarian cancer, pancreatic cancer, prostate cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonucleotide described in the present sequence is an immunostimulatory oligonulatory oligonulatory oligonulatory
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Pred. No. 9
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at risk of
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RESULT 886
ABL39403/c
ID ABL394
XX ABL394
XX ABL394
XX ABL394
XX Immuno
XX Immuno
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XX Synthe
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  RESULT 887
ABL39179/c
ID ABL391
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                                                                                                                                                                                                                  Query Match
Best Local S
Matches 17
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                                                                                                                                                                                                                                                                                                                                                                         The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, ossophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, mon-Hodgkin's lymphoma, melanoma, mysloma, oral cavity cancer, ovarian cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is an immunostimulatory oligonucleotide described in the
  ABL39179 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antibody-induced cell lysis; cancer; immunostimulatory; CD20; angiogenesis; metastasis; cytostatic; ss.
                                                                                                                                                                                                                                                                                                            Sequence 20 BP; 0 A; 0 C; 0 G; 20 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 309; 312pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Weiner G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       22-JUN-2001; 2001WO-US020154
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                                                                                                                                                                                                                                                                                                                                                             exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-JUN-2000; 2000US-0213346P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ABL39403
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                                                                                                                                                                   5393 AAAAAAATACAAAAAAGAAA 5412
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                                                                                                                          AAAAAAAAAAAAAAAA 1
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                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                                                                                                                                                                  Score 15.2; DB 1; Length Pred. No. 9.3e+02; o; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pred. No. 9.3e+02;
0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NO:
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RESULT 888
ABL54775/C
ID ABL547
XX ABL547
XX OABL547
XX OABL547
XX STOLL XX
DT 10-JUN
XX CD14 x
XX Single
XX Single
XX Single
XX Synthe
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to methods for treating or preventing cancer, involving administering to a subject having or at risk of developing cancer immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies. The methods are useful for treating or preventing cancer such as basal cell carcinoma, bladder cancer, bone cancer, brain and central nervous system (CNS) cancer, breast cancer, cervical cancer, colon and rectum cancer, connective tissue cancer, oesophageal cancer, eye cancer, kidney cancer, larynx cancer, leukaemia, liver cancer, lung cancer, Hodgkin's lymphoma, mon-Hodgkin's lymphoma, melanoma, myeloma, oral cavity cancer, ovarian cancer, stomach cancer, prostate cancer, rhabdomyosarcoma, skin cancer, stomach cancer, testicular cancer, and uterine cancer. The present sequence is a immunostimulatory oligonucleotide described in the
                                              Angiotensin-I converting enzyme; ACE; CD14; receptor; SNP; single-nucleotide polymorphism; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                                                                                                                                                                                                                    ABL54775;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Treating or preventing cancer, such as basal cell carcinoma, comprises administering immunostimulatory nucleic acids that induce expression of cell surface antigens and antibodies to a subject having or at risk of
                                                                                                                     CD14 receptor PCR primer SEQ ID NO 9.
                                                                                                                                                                      10-JUN-2002
                                                                                                                                                                                                                                                                   ABL54775 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         exemplification of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         developing cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (IOWA ) UNIV IOWA RES FOUND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Antibody-induced cell lysis; cancer; immunostimulatory; CD20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-APR-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                  5129
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                                                                                                                                                                                                                                                                                                                                                                                      AGGATCAGGAGCGACATGGA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP; 2 A; 8 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                                                                                      (first
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                                                                                                                                                                                                                                                                   DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.3%;
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                                                                                                                                                                      entry)
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Pred. No. 9.3e+02;
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The invention relates to a method for detecting 1 base polymorphism on the sequence of a chromosome or its fragment in which a sample nucleic acid is reacted with a reaction liquor containing a nucleic acid primer having a base adjacent to the polymorphic base at its 3'-end, one dideoxynuclectide corresponding to a polymorphic base having a distinguishable feature or its mixture, DNA polymerase and a composition required for its activity expression to detect the presence of taking dideoxynuclectide in the nucleic acid primer and to detect the type of the base to be specified. The method is used for detecting 1 base polymorphism on the sequence of a chromosome or its fragment. The present
                                                                                                                                                                  11-AUG-2000; 2000US-0224631P.
08-DEC-2000; 2000US-0254392P.
11-DEC-2000; 2000US-02535P.
12-JAN-2001; 2001US-00760500.
28-MAR-2001; 2001US-00820279.
                                                                                                                                                                                                                                                                                                                                                                                              07-MAR-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200218643-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nanoparticle-oligonucleotide; nanofabrication; nucleic acid detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nanoparticle-oligonucleotide #55
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                                                       Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABK65035 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      26-JUL-2000; 2000JP-00225354
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                                                                                                                 (NANO-) NANOSPHERE INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          is that of a PCR primer, useful
                                  Letsinger RL, Garimella V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page 10; 10pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
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                                  Ŀ
                             Mucic RC,
i Z, Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0;
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Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                          Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in examples of the invention
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                                                          Elghanian
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CC to the sequence of NA; (b) contacting NA and NP under conditions
CC effective to allow hybridisation of OGN with NA; and (c) observing a
CC detectable change brought about by hybridisation of OGN with NA. The
CC method is useful for detecting a nucleic acid, separating a selected
CC nucleic acid from others and methods of nanofabrication. Detecting
CC analytes such as nucleic acids and proteins are useful for the diagnosis
CC of genetic, bacterial and viral diseases. The OGN-NP conjugates that use
CC cyclic disulphide linkers improve the sensitivity of diagnostic assays.
CC in particular assays using OGN-NP conjugates prepared using linkers
CC comprising a steroid residue attached to a cyclic disulphide have been
CC conjugates prepared using alkanethiols or acyclic disulphides as the
CC conjugates prepared using alkanethiols or acyclic disulphides as the
CC conjugates prepared using alkanethiols or acyclic disulphides as the
CC conjugates prepared using alkanethiols or acyclic disulphides to a DNA
CC target to be PCR amplified can be carried through the 30 or 40 heating
CC conjug cycles of the PCR and are still able to detect the amplicons
CC without opening the tubes and causing contamination. ABK64981-ABK65055
CC represent nanoparticle-oligonucleotides of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a method of detecting a nucleic acid (NA) having at least 2 portions comprising: (a) providing nanoparticles (NP) with attached oligonuclectides (OGN), where OGN has a sequence complementary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Detecting nucleic acid, useful for diagnosis of genetic, vin bacterial disease, comprises hybridizing nanoparticles with oligonucleotides to nucleic acid and detecting change brough
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 18; Page 410; 412pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             change brought about by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    acid (NA) having
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 attached
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유 S Matches Query Match Local 5393 AAAAAAATACAAAAAAGAAA 5412 17; Conserv -AAAAAAAAAAAAAAAAAA 20 Conservative 0.3%; <u>.</u> Score 15.2; DB 1 Pred. No. 9.3e+02 Mismatcheв DB 1; Length 20 Indels <u>,</u> 0

Sequence

20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;

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RESULT 890
ABK65050
ID ABK650
XX ABK650
XX ABK650
XX Nanopa
XX Nanopa
XX Nanopa
XX Nanopa
XX NANOpa
XX NANOpa
XX NO2002
XX WO2002
XX HO2002
XX HO2
                                                                                                                                                                                                                                                            11-AUG-2000; 2000US-0224631P.
08-DEC-2000; 2000US-0254392P.
11-DEC-2000; 2000US-0255235P.
12-UAN-2001; 2001US-00760500.
28-MAR-2001; 2001US-00820279.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nanoparticle-oligonucleotide; nanofabrication; nucleic acid detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nanoparticle-oligonucleotide #70.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABK65050 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-AUG-2001; 2001WO-US025237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO200218643-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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Mirkin CA, Letsinger RL, Muc Taton TA, Garimella V, Li Z,

Mucic RC, S i Z, Park S;

Storhoff JJ,

Elghanian

(NANO-) NANOSPHERE INC.

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RESULT 891
ABS52449
ID ABS524
XX ABS524
XX ABS524
XX IS-NOV
DT 15-NOV
XX ISS; CKW ORP2;
KW System
KW INSULI
KW INSULI
KW Expenial
KW Expenial
KW Expenial
KW Expenial
KW INSULI

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC effective to allow hybridisation of OGN with NA, and (c) observing a CC detectable change brought about by hybridisation of OGN with NA. The method is useful for detecting a nucleic acid, separating a selected cc mucleic acid from others and methods of nanofabrication. Detecting canalytes such as nucleic acids and proteins are useful for the diagnosis of genetic, bacterial and viral diseases. The OGN-NP conjugates that use cyclic disulphide linkers improve the sensitivity of diagnostic assays. CI in particular assays using OGN-NP conjugates prepared using linkers comprising a steroid residue attached to a cyclic disulphide have been conjugates prepared using alkanethiols or acyclic disulphides as the CI inker. The OGN-NP conjugates are stable allowing them to be used circuity in PCR solutions. Therefore conjugates added as probes to a DNA carget to be PCR amplified can be carried through the 30 or 40 heating cooling cycles of the PCR and are still able to detect the amplicons without opening the tubes and causing contamination. ABK64981-ABK65055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                       ss; long interspersed nuclear element; LINE-1; p40; PCR; primer; ORF1; ORF2; L1; Alzheimer's disease; autoimmune disease; schizophrenia; systemic lupus erythematosus; multiple sclerosis; scleroderma; insulin-dependent diabetes mellitus; rheumatoid arthritis; phemphigus; psoriasis; autoimmune thyroid disease; polymyositis; vitiligo; mixed connective tissue disease; dermatomyositis; Sjogren's syndrome; pemphigoid; primary biliary cirrhosis; chronic active hepatitis; Crohn's disease; ulcerative colitis; pernicious anaemia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to a method of detecting a nucleic acid (NA) havin at least 2 portions comprising: (a) providing nanoparticles (NP) with attached oligonucleotides (OGN), where OGN has a sequence complementary to the sequence of NA; (b) contacting NA and NP under conditions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          oligonucleotides to nucleic acid and detecting change brought about by hybridization.
                                                                         19-DEC-2000; 2000US-0256673P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human LINE-1 DNA associated PCR primer #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-NOV-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABS52459 standard; DNA; 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 24; Fig 44; 412pp; English.
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                                                                                                                                                                                                                                                                                          WO200262197-A2
                                                                                                                                               19-DEC-2001; 2001WO-US049353
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 АААААААААААААААААААА
   HOSPITAL FOR SPECIAL SURGERY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 15.2;
Pred. No. 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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RASULT 892
AAL45122/c
ID AAL451
XX
AC AAL451
XX
DT 24-MAY
XX
OI 1900
XX
CN Oligon
KW Oligon
KW Oxidat
XX
XX
OS Synthe
XX

24-MAY-2002

(first entry

AAL45122

AAL45122 standard; DNA; 20

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0,

Oligonucleotide synthesis method related DNA #1

Oligonucleotide synthesis; polynucleotide array; protecting group;

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CC (Li=long interspersed nuclear element, LINE-1). Also included are (1) CC identifying an individual at risk for or suffering from a complex disease CC comprises: (a) identifying or detecting the amount of intronic regions of genes containing full length Li elements or in 5' or 3' regulatory CC regions of genes containing a full length high fidelity consensus L1 CS sequence of the individual's DNA from a sample; and (b) comparing the presence of the Li sequence or its amount in the intronic regions of CC genes or the 5' or 3' regulatory regions with a control sample of DNA CC form an individual not susceptible to or at risk for or currently cCC suffering from a complex disease, where the genes identified are involved in a complex genome; (2) treating or preventing a complex disease by CC antibody directed against Li mRNA, and an antibody directed against in mRNA or protein particles having Li risk for or suffering from a complex disease, by detecting antibodies or auto CC antibodies directed against to mRNA or protein products which indicates that the CC individual is at risk for or suffering from a complex disease or auto CC antibodies directed against ribonucleo-protein particles having Li mRNA CC individual is at risk for or suffering from a complex disease or auto CC elements, disease, autoimmune diseases, schizophrenia, systemic lupus crythematosus, multiple sclerosis, insulin-dependent diabetes mellitus, complexis and particlus and permicious anaemia). Detection of the protein products of Li elements, either ORF1/P40 or ORF2 gene products of be used to indicate the present can induce or exacerbate autoimmune disease. The present cc colitis and permicious anaemia). Detection of the protein products of Li elements, either ORF1/P40 or ORF2 gene products can be used to indicate the present can induce or exacerbate autoimmune disease. The present can induce or exacerbate autoimmune disease.
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                                       S
                                                                                         Query Match
Best Local S
Matches 17
                                                                                                                                                                                       Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 comprising identifying genes containing full-length L1 elements in their intronic region or containing a full length L1 element with high sequence fidelity to the L1 consensus sequence in their 5 or 3 regulatory region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 137; 138pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Identifying a gene involved in a complex disease, e.g. schizophrenia, comprises detecting genes having full-length L1 element in their intronic region or high sequence fidelity to L1 consensus sequence in the 5' or 3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-643381/69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to identifying a gene involved in a complex disease
                                               3590 ATGTTGCTCAGGCTAATCTC 3609
                                                                                                                    Similarity
                                                                                                                                                                                            20
ATGTTGGCCAGGCTGATCTC 20
                                                                                                                                                                                    BP; 3 A; 5 C; 6 G; 6 T; 0
                                                                                              Conservative
                                                                                                                 0.3%;
                                                                                              0
                                                                                                                    Score 15.2;
Pred. No. 9
                                                                                              Mismatches
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. 9.3e+02;
3;
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                                                                                                                                                                                            0 Other;
                                                                                                                                          1; Length 20;
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                                                                                              Gaps
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RESULT 893
ABQ80721
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to a method for the synthesis of a polynucleotide which involves coupling a second nucleoside to a first nucleoside through a phosphite linkage, where the second nucleoside has a non-carbonate protecting group protecting a hydroxyl, and exposing the product to a composition which concurrently oxidizes the phosphite formed to a phosphate and deprotects the protected hydroxyl of the second nucleoside. The method to second for synthesizing the polynucleotides, for carrying out either 3' to 5' or 5' to 3' synthesis and for fabricating an addressable array of polynucleotides on a substrate. The present sequence is an oligonucleotide produced to demonstrate the method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthesis of polynucleotide useful during fabrication of an array involves coupling nucleoside phosphoramidite and a solid-supported nucleoside and treating the product with an oxidation/deprotection
                                                                                                                                                                                                Salmonella
                                                                                                                                                                                                                                             Salmonella toxin
Novel oligonucleotide for detecting Salmonella toxin gene inva or
                                                                                                17-JAN-2001; 2001JP-00009464
                                                                                                                       17-JAN-2002; 2002EP-00001254
                                                                                                                                                                       EP1233073-A2
                                                                                                                                                                                                                                                                       22-NOV-2002
                                                                                                                                                                                                                                                                                              ABQ80721;
                                                                                                                                                                                                                                                                                                                      ABQ80721 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         involves coupling nucleonucleoside and treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   28-JUL-2000; 2000US-00627249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-JUL-2001; 2001EP-00118360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30-JAN-2002.
                         WPI; 2002-668421/72.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dellinger DJ,
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                                                                        ( LYOT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (AGIL-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         the invention
                                                                                                                                                                                                                                                                                                                                                                                                          5393 AAAAAAATACAAAAAAGAAA 5412
                                                                                                                                                                                                                                                                                                                                                                                  20
                                                                                                                                                                                                                     invA; stn;
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                                                                         TOSOH
                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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                                                                                                                                                                                                                                                                                                                                                                                   AAAAAAAAAAAAAAAAAAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BP; 0 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                Ishiguro
                                                                                                                                                                                                                                                                      (first
                                                                         CORP
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                                                                                                                                                                                                                                             gene stn mRNA-binding oligonucleotide #6.
                                                                                                                                                                                                                     primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     36pp;
                                                                                                                                                                                                                                                                      entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       English
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.2;
Pred. No. 9.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 T; 0 U;
                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Caruthers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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RESULT 85
ABL36232
AC ABL3
AC 
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                                                    The present invention relates to a method of inhibiting skin inflammation associated with a skin disorder selected from psoriasis, atopic dermatitis and allergic contact dermatitis, which involves administering a composition containing delipidated and deglycolipidated Mycobacterium vaccae cells or M. vaccae culture filtrate. The skin disorder to be treated may also include alopecia areata, and skin cancers such as basal cell carcinoma, squamous cell carcinoma and melanoma. The composition acts by inhibiting the Th2 immune response. The present sequence is a probe described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Skin disorder; psoriasis; atopic dermatitis; allergic contact alopecia areata; skin cancer; Mycobacterium vaccae; melanoma; antipsoriatic; dermatological; antiinflammatory; antiallergic;
                                                                                                                                                                                                                                                                                                                                                                                                       Inhibiting skin inflammation associated with skin disorder e.g. psoriasis, by administering composition comprising delipidated deglycolipidated Mycobacterium vaccae cells or Mycobacterium va
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Pred. No. 9.3e+02;
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                                               cells in which both alleles of a gene are modified, comprising modifying one allele by insertion or replacement by a cassette having an ceremon compressible selectable marker and modifying other allele by crecombination, of a promoter replacement fragment with a heterologous promoter, so that expression of the second allele is regulated by the promoter. (M1) is useful for constructing a strain of diploid fungal cells in which both alleles of a gene are modified. The diploid fungal cells having both alleles modified are useful for identifying a gene that cells having both alleles modified are useful for identifying a gene that contributes to the virulence and/or pathogenicity of a fungus, a gene that contributes to the resistance of a diploid fungus to an antifungal agent that inhibits the growth of a diploid fungus and for identifying a therapeutic agent for treatment of a mammalian compound catabolism, biosynthetic, transporter, transcriptional, compound catabolism, biosynthetic, transporter, transcriptional, resulting infection by C. albicans. The present sequence is that of a PCR primer used in the method of the invention. Note: The sequence data for this patent is not represented in the printed specification batt is based on information suntied to Dervent by the Broncen patent offices.
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Best Local Similarity
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                                        on sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to constructing (M1) a strain of diploid fungal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 36; SEQ ID NO 4750; 167pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Constructing strains for identifying gene products as effective targets for therapeutic intervention, by inaccivating in the strain one allele of a gene and placing other allele of the gene under conditional expression.
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20-FEB-2001; 2001US-00792024
22-AUG-2001; 2001US-0314050P
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                                        information supplied to Derwent by the European Patent Office
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Pred. No. 9
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9.3e+02;
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                                                                                  The invention relates to bone strength and mineralisation regulatory correction (BSWR) and its corresponding nucleic acid sequence. BSWR DNA is useful for the diagnosis or therapy of osteoporosis and for regulating (increasing) bone strength and mineralisation in a human subject by activating a bone density regulating transmembrane receptor (BSWR DIA). An expression vector comprising a promoter that is operably linked to BSWR DNA is useful for modulating bone density and for enhancing bone strength and mineralisation in a mammal cell. Composition comprising a BSWR effector is useful for treating osteoporosis and is useful particularly as a coating for prosthetic devices and surgical implants. BSWR is useful for screening lead pharmaceutical agents as BSWR effectors, which may be used to treat a range of eye disorders such as caseful particularly, hypertensive retinopathy and retinopathy of cular vessels is disrupted. The present sequence is a PCR primer used to amplify cDNA and gDNA molecules useful for detecting polymorphic BSWR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 20 BP; 2 A; 5 C; 6 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Polynucleotide encoding bone protein useful for diagnosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-AUG-2000; 2000US-0226119P.
22-SEP-2000; 2000US-0234337P.
13-JUL-2001; 2001US-0304851P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        bone strength; mineralisation; ophthalmological; antidiabetic; bone density regulating transmembrane receptor; prosthetic device; surgical implant; diabetic retinopathy; hypertensive retinopathy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAD34368 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Fig 5; 124pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (AVET )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-AUG-2001; 2001WO-US041788
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          therapy; osteoporosis; prematurity; ocular vessel; eye disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; bone strength and mineralisation regulatory protein; BSMR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human BSMR gene polymorphism detecting PCR primer, LRGEN23R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-JUL-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5070 TCATCTGGTGGCCACAGCAG 5089
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85.0%; Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Olsen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       strength and mineralization regulatory or therapy of osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rawadi G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     9.3e+02
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Sequence 20

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                        two portions, involving providing nanoparticles having oligonucleotides attached to it, which has a sequence complementary to sequence of two portions of nucleic acid, contacting nucleic acid and nanoparticles, to allow hybridisation of oligonucleotides with two or more portions of nucleic acid, and observing a detectable change brought about by hybridisation. (M1), nanoparticles (I), nanoparticle-oligonucleotide conjugates (II) and the aggregate probe are useful for detecting two or more nucleic acids (from a biological source) having at least two or more nucleic acids (from a biological source) having at least two or portions, such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic, or structurally-modified natural or synthetic RNA or DNA, or a product of a polymerase chain reaction amplification.

(II) is useful for preparing a nunceptobe conjugate for detecting an analyte, and for detecting a nucleic acid bound to an electrode surface.

(I) and (II) are useful for fabrication, and for separating a selected nucleic acid having two portions from other nucleic acids. (I), (II) and the aggregate probe are useful for detecting an analyte (especially polyvalent analyte) in a sample. This sequence represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and nanoparticles to allow hybridization, and observing detectable change.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mirkin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 18; Page 437; 442pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2002-608256/65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-DEC-2000;
11-DEC-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid detection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention describes a method of detecting (M1) a nucleic acid having
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NANO-) NANOSPHERE INC.
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Garimella V, Li
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; 2001US-00820279.
; 2001US-0282640P.
; 2001US-00927777.
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2000US-0255235P.
2000US-0255236P.
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Pred. No. 9
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Matches 17
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11-DEC-2000; 2000US-0255236P.
12-JAN-2001; 2001US-00760500.
28-MAR-2001; 2001US-0082079.
09-APR-2001; 2001US-0282640P.
10-AUG-2001; 2001US-00927777.
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nucleic acid, and observing a detectable change brought about by hybridisation. (M1), nanoparticles (I), nanoparticle-oligonucleotide conjugates (II) and the aggregate probe are useful for detecting two or more nucleic acids (from a biological source) having at least two portions, such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic, or structurally-modified natural or synthetic RNA or DNA, or a product of a polymerase chain reaction amplification. (II) is useful for repearing a nanoprobe conjugate for detecting an analyte, and for detecting a nucleic acid bound to an electrode surface. (I) and (II) are useful for fabrication, and for separating a selected nucleic acid having two portions from other nucleic acids. (I), (II) and the aggregate probe are useful for detecting an analyte (especially polyvalent analyte) in a sample. This sequence represents a
                                                                                                                                                                                                                                                                                                                                                                                                                                   two portions, involving providing nanoparticles having oligonucleotides attached to it, which has a sequence complementary to sequence of two portions of nucleic acid, contacting nucleic acid and nanoparticles, to allow hybridisation of oligonucleotides with two or more portions of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and nanoparticles to allow hybridization, and observing detectable change.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-DEC-2000; 2000US-0254392P

08-DEC-2000; 2000US-0254418P

11-DEC-2000; 2000US-0255235P

11-DEC-2000; 2000US-0255236P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 24; Fig 44; 442pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2002-608256/65.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-DEC-2001; 2001WO-US046418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          13-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WO200246472-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention describes a method of detecting (M1) a nucleic acid having
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NANO-) NANOSPHERE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5393 AAAAAAATACAAAAAAGAAA 5412
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Garimella V, Li Z,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
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Pred. No. 9.3e+02;
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RESULT 900
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ABN86365/c
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                                                                                                                        Query Match
Best Local S
Matches 17
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Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                    The invention relates to the treatment of paramyxovirus infection that involves administration of an active ingredient such as RANTES chemokine (I) or an expression system (II). (I) and (II) are used in the treatment and diagnosis of paramyxoviral infection, especially in children. RANTES chemokine has a significant effect on respiratory infections caused by paramyxovirus. RANTES acts downstream of viral entry and signals through specific CCR1 and/or CCR5 chemokine receptors to interrupt the death pathway of macrophages which have been infected by virus. RANTES not only inhibits apoptosis of infected macrophage but also clears the macrophage of infection. The present sequence represents a PCR primer specific for
                                                                                                                                                                               Sequence 20 BP; 5 A; 0 C; 11 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                    Use of RANTES chemokine or an expression system in treatment and diagnosis of paramyxovirus infection, especially in children.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-OCT-2000; 2000US-0243264P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     18-JUL-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Paramyxovirus; infection; apoptosis; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RANTES DNA amplifying primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABN86365;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABN86365 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     polynucleotide used to demonstrate the method of the
                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 8; 19pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        23-OCT-2001; 2001WO-US045244
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                                                                                              228 CCCTCACCCTCACCCTCCCT 247
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                                                                                                                                        Similarity
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  standard; DNA;
                                                                      CCCTCACCATCATCCTCACT 1
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                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  infection; RANTES; chemokine; antiviral; CCR1; CCR5;
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                                                                                                                         0; Mismatches
                                                                                                                                     Pred.
                                                                                                                                      Score 15.2;
Pred. No. 9.
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Pred. No. 9.3e+02;
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                                                                                                                                      DB 1; Length 20; .3e+02;
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ABI96122/
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                                                                                          RESULT 901
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Best Local S
Matches 17
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07-JUL-1997;
06-JUL-1998;
                                                                                                                                                                                                                                                                         The present invention relates to novel antisense oligonucleotides which are targetted to nucleic acids encoding human raf proteins and capable cinhibiting raf expression. The invention also relates to methods of inhibiting hyperproliferation of cells which involves contacting the hyperproliferating cells with a therapeutically effective amount of an oligonucleotide of the invention. The method is useful for treating cancer, angiogenesis or neovascularisation, especially ocular angiogenesis or neovascularisation. The present DNA sequence is an angiogenesis or neovascularisation. The present DNA sequence is an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human; raf; hyperproliferation; neovascularisation; ocular angiogenesis; therapy; cancer; cytostatic; anti-angiogenic; vascular; ophthalmological; antisense; phosphorothicate backbone; B-raf kinase; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human B-raf kinase antisense
                                                                                                                                                                                                                                      Sequence 20 BP; 2 A; 3 C; 1 G; 14 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-DEC-2002
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                                       ABI96122;
                                                                                                                                                                                                                                                                                                                                                                                                                       Treating cancer, angiogenesis or neovascularization by administering antisense oligonucleotides targeted to human raf sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Monia
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       modified_base
                                                                 ABI96122 standard;
                                                                                                                                                                                                                                                              antisense oligonucleotide targetted to human B-raf kinase
                                                                                                                                                                                                                                                                                                                                                                                                Example 18; Col 26; 41pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (ISIS-) ISIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31-MAY-1995;
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                                                                                                                                                         5412 AAAATGAAAATAAAGGAATA 5431
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ilarity 85.0%;
Conservative (
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96US-00756806.
97US-00888982.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  note= "Phosphorothioate backbone"
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                                                                 DNA;
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                                                                 BP
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Pred. No. 9.
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                                                                                                                                                                                                            DB 1; Length 20;
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16-FEB-2002

(first entry)

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RESULT 902
ABN86937/c
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AC ABN869
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DT 29-JUL
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                                                                                                                                                                                                                                                                                                                                                                                                    CC oligonuclectide probes (II) will hybridise with little mismatch, where CC (I) have melting temperatures within a narrow range. The method is useful CC for detecting infectious diseases caused by bacterial infectious agents ce.g. Salmonella, Listeria monocytogenes and Haemophilus influenza, fungal CC infectious agents e.g. Cryptococcus necformans, Candida albicans and CC Aspergillus fundyautus, viruses e.g. T-cell lymphocytotrophis cirus, CC Epstein-Barr virus and polio virus, and parasitic infectious agents celected from Onchoverva volvulus, Entamoeba histolytica and Dracunculus celected from Marconcolus, Entamoeba histolytica and Dracunculus celected from Sancer involving oncogenes, tumour suppressor genes, or genes cinvolved in DNA amplification, replication, recombination or repair, the CC cancer is specifically associated with a gene selected from BRCA1 gene, CC method is also used for environmental monitoring, forensics and the food and feed industry, detecting comprises scanning (using e.g. a scanning celectron microscope and infrared microscope) the support at the CC presence or absence of the target nucleotide sequences. AB182074 to a fermination of the oligonucleotide probe celected from to aligonucleotide sequences used in the exemplification of the neasent loligonucleotide sequences used in the exemplification.
                                                                                                                                                                                                                                                                       Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; K-ras; PCR primer; probe; capture probe; mutation detection; ligase detection reaction; LDR; p53; BRCA1; BRCA1; infectious disease; infection; 21 hydroxylase deficiency; Turner Syndacome; obesity; cancer; oncogene; tumour suppressor; human papillomavirus; forensic; environmental monitoring; food industry; feed industry; ss.
                                                                                                                                                                                                                                                                                                                                                      Sequence 20
                      29-JUL-2002
                                                            ABN86937;
                                                                                                 ABN86937 etandard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Designing capture oligonucleotide probes complementary oligonucleotides hybridize
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-034366/04.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention describes a method (M1) for designing capture
                                                                                                                                                                                                                                  3392 GGCTGGACCAGACAACTCTG 3411
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                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5; Fig 29; 300pp; English.
                                                                                                                                                                                                  GGCTGGTCCAGACAAGGCTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Zirvi M,
                                                                                                                                                                                                                                                                                                                                                      BP; 3
                                                                                                                                                                                                                                                                           Conservative
                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                            invention
                                                                                                                                                                                                                                                                                                                                                      A; 8 C; 5 G; 4 T; 0
                                                                                               DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FOUND INC.
                                                                                                                                                                                                                                                                                           0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gerry NP,
                                                                                                                                                                                                                                                                         0;
                                                                                                   ₽P
                                                                                                                                                                                                                                                                                                             Score 15.2;
                                                                                                                                                                                                                                                                                               Pred.
                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Favis
                                                                                                                                                                                                                                                                                               No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ¤
                                                                                                                                                                                                                                                                                                                                                      U; 0 Other;
                                                                                                                                                                                                                                                                                           9.3e+02
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for use on a support to which with little mismatch.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kliman
                                                                                                                                                                                                                                                                                                             DB 1; Length 20
                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                           0
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Grosse WM,
Kekuda R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; NOVX; cytostatic; antiarteriosclerotic; cardiovascular; lymphoma; antidiabetic; immunosuppressive; neuroprotective; gene therapy; cancer; cardiomyopathy; atherosclerosis; cell signal processing; diabetes; AIDS; metabolic pathway modulation; neoplastic; neurological disorder; asthma; adenocarcinoma; prostate cancer; uterus cancer; immune response; Crohn's disease; multiple sclerosis; Graft versus host disease;
                                                                                                                                                                                                                                                                                                                                                                                     16-OCT-2000;
16-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                             16-OCT-2000; 2000US-0240648P.
16-OCT-2000; 2000US-0240662P.
16-OCT-2000; 2000US-0240669P.
16-OCT-2000; 2000US-0240703P.
                                                                                                                                                                                                                                                                                                                                                                                                                                               12-OCT-2000; 2000US-0240113P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12-OCT-2001; 2001WO-US031922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200230974-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PCR primer; 88.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human NOV4 reverse PCR primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                              16-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                     18-JAN-2001;
                                                                                                                                                                                                                                                                                                                                                   (CURA-)
                                                                                                                                                                                                                                                                                                                                           CURAGEN CORP.
                                                                                                                                                                                                                                                                                                                                                                   2000US-0240732P.
2000US-0241190P.
2001US-0262455P.
                                                                                                                                                                                                                                                                                                                                                                                                                               2000US-0240637P
                                                                                                                                                                                                                                                                                                                                                                                                                                        2000US-0240625P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NO:56
                                                                                                                                                                                                                                                                                                                          Burgess CE,
                                                                                                                                                                                                                                                                                                                           Mishra V;
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New NOVX polypeptides and polynucleotides, useful for treating or preventing a NOVX-associated disorder or a pathological state in a subject, particularly a human, e.g. cardiomyopathy, atherosclerosis, Edinger S, 2002-444172/47. Li L, Pa , Gerlach Alsobrook JP, Lepley DM, Burg ۲, Macdougall J, Stone D, Zerhusen BD, D, Gunther E, Spytek KA; , Ellerman

Example 2; Page 178; 227pp; English.

cancer or diabetes.

The present invention describes novel human proteins designated NOVX CC (where X is 1, 2a, 2b, 2c, 2d, 3, 4, 5, 6a, 6b, 7, 8, or 9). NOVI is a CC tyrosine-protein kinase 6-like protein; NOV2a-d are keratin 4-like CC protein; NOV3 is a collagen-like protein; NOV4 is a cystatin B-like protein; NOV8 is a serotonin receptor-like protein; NOV6a and NOV55v are CC cold inducible glycoprotein 30-like protein; NOV7 is a matrilin-2-like protein; NOV8 is a leukocyte surface antigen (CD53)-like protein; and CC NOV9 is a tyrosine kinase-like protein. NOVX sequences have cytostatic, CC antiarteriosclerotic, cardiovascular, antidiabetic, immunosuppressive and CC neuroprotective activities, and can be used in gene therapy. The NOVX cc sequences can be used in therapeutics, particularly for treating, CC preventing or alleviating a NOVX-associtated disorder or a pathological ctate in a subject, particularly a human. These disorders include cardiomyopathy, atherosclerosis, a disorder related to cell signal CC processing and metabolic pathway modulation or diabetes. The NOVX csequences are also useful for determining the presence of or predisposition to a disease associated with altered levels of NOVX caplopetide or nucleic acid, particularly cancer. The NOVX sequences are copolastic or neurological disorders, and in the treatment of caplositions, lymphoma, prostate cancer, uterus cancer, immune CC response, AIDS, asthma, Crohn's disease, multiple sclerosis or Graft versus host disease. The present sequence represents a PCR primer for human NOV4, which is used in an example from the present invention

Sequence 20 BP; 3 A; 3 C; 6 G; 8 T; 0 U; 0 Other;

Score 15.2;

DB 1;

Length 20;

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ARBSULT 903
ABN87103/c
ID ABN8711
AC ABN871
XX ABN871
XX Protei
XX Protei
XX Synthe
XX Synthe
XX Synthe
XX Ho-OC7
XX 16-OC7
XX 16-OC7
XX (PHYL-
XX New nx
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XX New nx
PT reseal
PT design
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                                                                                                                                                                                       CC protein having a mutated amino acid sequence, where the non-antibody corporein having a mutated amino acid sequence, where the non-antibody corporein hinds with a Kd at least as tight as 10 nM to a compound that is come to bound as tightly by the reference protein. The non-antibody protein is useful as scaffolds for selecting or designing a protein framework continues the properties, e.g. for binding any antigen of cinterest, or for destroying or inactivating antibody molecules. The non-antibody protein is also useful in all areas where antibodies are used, compared the respective or diagnostic fields, and for screening novel binding proteins useful in the above-mentioned fields. The present compounds the properties superior to those of natural compounds and can be evolved rapidly in vitro. The present proteins or cantibodies, and can be evolved rapidly in vitro. The present proteins or cantibody mimics exhibit improved biophysical properties, such as concentrations. In addition, these molecules are readily expressed and concentrations. In addition, these molecules are readily expressed and concentrations. In addition, these molecules are readily expressed and concentrations. In addition, these molecules are readily expressed and concentration yeate systems (e.g. Fabcherichia coll), in eukaryotic systems (e.g. rabbit concentration end to the present sequence is used in the sequence of concentration, e.g. in vitro selection using RNA-protein fusion technology, phage display or yeast display systems. The present sequence is used in the exemplification of the present invention
          Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 85.0 Matches 17; Conservative
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                                                                                                                                            Sequence 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 58; 94pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New non-antibody proteins having an immunoglobulin fold, useful in research, therapeutic or diagnostic fields, particularly as scaffo designing proteins with specific properties, e.g. for binding any
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-444238/47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16-OCT-2000; 2000US-00688566
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      25-APR-2002
Local 5.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PHYL-) PHYLOS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               present invention describes a non-antibody protein, comprising ain having an immunoglobulin-like fold, derived from a reference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4816
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                                           Similarity
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                                                                                                                                        BP;
                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
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                                                                                                                                        ₽,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RW,
                                        0.3%;
                                                                                                                                        0 C; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            85.0%;
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0;
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                                                                                                                                        G; 20 T; 0 U; 0 Other;
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          Score 15.2; DB 1; Length 20; Pred. No. 9.3e+02; O; Mismatches 3; Indels
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          0
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          Gaps
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RESULT 904
AAD41716/c
ID AAD417
XX AAD417
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XX Human
XX Human
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XX Homo (
OS Synth(
XX Key
FT modif:
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                                                                                                                                             Novel antisense compounds targeted to nucleic acids encoding interleukin-
12 p35 subunit, useful for modulating interleukin-12 p35 subunit expression and treating diseases associated with expression of the subunit in humans.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                    07-MAY-2001; 2001US-00851520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07-MAY-2001; 2001US-00851520.
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                                                                                                  Claim 3;
                                                                                                                                                                                                                                                                                                                       Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             modified_base
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                                                                                                                                                                                                                                                                                                                                                                       (ISIS-) ISIS PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5393 ААААААТАСАААААСААА 5412
                                                                                                                                                                                                                                                                         2002-535980/57.
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                                                                                                Col 49-50; 44pp; English
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                                                                                                                                                                                                                                                                                                                       Freier SM;
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/mod_base= r
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16. .20
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/mod_ba
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/mod_ba
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/note= "2'-methoxyethyl (MOE) residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mod_base=
12. .13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note= "2'-methoxyethyl (MOE) residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         note= "Phosphorothicate backbone"
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The present invention relates to novel antisense oligonucleotides which specifically hybridise with specific regions of nucleic acids encoding interleukin-12 (IL-12) p35 subunit and inhibit the expression of human 1

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RESULT 905
ACA63945/c
ID ACA63945 standard; D:
XX
AC ACA63945;
XX
AC ACA63945;
XX
DT 16-JUN-2003 (first
DT 16-JUN-2003)
Cfirst
XX
DZ Novel human secreted
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DZ Novel human secreted
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DZ Novel human secreted
XX
Human; secreted and
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Human; secreted and
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XX
US2002192706-A1.
XX
US2002192706-A1.
XX
DD 19-DEC-2001; 2001US-
XX
PD 19-DEC-2001; 27US-
PR 24-OCT-2001; 27US-
PR 13-NOV-1997; 97US-
PR 13-NOV-1997; 97US-
PR 13-NOV-1997; 97US-
PR 11-MAR-1998; 98US-
PR 12-MAR-1998; 98US-
PR 20-MAR-1998; 98US-
PR 20
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              -12 p35 subunit. Sequences of the invention are useful for inhibiting the expression of human IL-12 p35 subunit in human cells or tissues and for treating animals, particularly humans suspected of having or being prone to diseases or conditions associated with expression of IL-12 p35 subunit. They are useful for diagnostics, therapeutics and as research reagent, e.g. prophylactically to prevent or delay infection, tumour formation or inflammation. Sequences of the invention are useful for antisense therapy. The present sequence is an antisense oligonucleotide targetted to human IL-12 p35 subunit DNA. This sequence is used in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human; secreted and transmembrane protein; PRO; antiinflammatory; antiarteriosclerotic; cardiant; anti-infertility; anti-HIV; cytostatic; antidiabetic; gene thorapy; inflammatory disease; organ failure; atherosclerosis; cardiac injury; inflammatory the defect; premature aging; AIDS; cancer; diabetic complication; chromosome mapping gene mapping; pharmaceutical; diagnostic; biosensor; bioreactor; tissue typing; PCR; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 20
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l Similarity 85.0%;
17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      secreted and transmembrane protein related primer #237.
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      97US-0062250P

97US-0064249P

97US-0065311P

98US-0077632P

98US-0077641P

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98US-0077641P

98US-0077804P

98US-0078004P

98US-0078930P

98US-0078930P

98US-0078939P

98US-0079294P

98US-0079294P

98US-00792664P

98US-0079664P

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31-MAR-1998

01-APR-1998

01-APR-1998

08-APR-1998

08-APR-1998

09-APR-1998

15-APR-1998

15-APR-1998

15-APR-1998

15-APR-1998

15-APR-1998

15-APR-1998

15-APR-1998

15-APR-1998

15-APR-1998

22-APR-1998

22-APR-1998

22-APR-1998

22-APR-1998

22-APR-1998

23-APR-1998

23-APR-1998

24-APR-1998

25-APR-1998

27-APR-1998

27-APR-1998

28-APR-1998

29-APR-1998

29-APR-1999

30-DEC-1999

30-DE
                                                  Ashkenazi AJ
Ferrara N,
Goddard A,
Kljavin IJ,
Stewart TA,
           WPI; 2003-328860/31.
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                                               zi AJ, Baker KP, Botstein D,
N, Filvaroff B, Fong S, Gg
A, Godowski PJ, Grimaldi JC
IJ, Kuo SS, Napier MA, Par
TA, Tumas D, Williams PM,
                                                                                                                                                                                      GENENTECH INC.
                                                                                                                                                                                                                             99WO-US03124

99WO-US03124

99WO-US031274

2000WO-US0302219

2000WO-US000277

2000WO-US000376

2000WO-US004341

2000WO-US005841

2000WO-US006319

2000WO-US006319

2000WO-US00753

2000WO-US013705

2000WO-US013705

2000WO-US013705

2000WO-US013705

2000WO-US013705

2000WO-US013705

2000WO-US013705

2000WO-US013705

2001WO-US03496

2001WO-US03496

2001WO-US01705

2001WO-US01705

2001WO-US01705

2001WO-US01705

2001WO-US01706

2001WO-US01706
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98US-0080132P

98US-0080332P

98US-0080333P

98US-0081070P

98US-0081071P

98US-0081203P

98US-0081203P

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98US-0081855P

98US-0081855P

98US-0081855P

98US-0082704P

98US-0082704P

98US-008271111

98WO-US01021855

99WO-US010733

99WO-US010733

99WO-US01255

99WO-US01255

99WO-US01255

99WO-US01255

99WO-US01255
                                                    Botstein D, Desnoyers L, Eaton DL;

Pong S, Gao W, Gerber H, Gerritsen ME;

Grimaldi JC, Gurney AL, Hillan KJ;

ier MA, Pan J, Paoni NF, Roy MA, Shelt-

liams PM, Wood WI;
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Shelton

DL;

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RESULT 906
ADA4471/C
ID ADA447
XX ADA447
XX ADA447
XX Antise
XW Antise
XW Antise
XW Antise
XW Antise
XW Antise
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention describes an isolated nucleic acid (I) comprising, or which is at least 80 % sequence identity to, or the full-length coding sequence of, any of 118 300-2100 nucleotide sequences, which encodes its corresponding PRO polypeptide selected from 118 100-700 amino acid sequences, all given in the specification. The nucleic acids and polypeptides are useful for treating inflammatory diseases, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, cancer, or diabetic complications. The nucleic acids are useful as hybridiastion probes, in chromosome and gene mapping, and in generating antisense RNA or DNA. The polypeptides are useful as pharmacceuticals, diagnostics, biosensors or bioreactors. Both are useful in tissue typing. This sequence represents a novel human secreted and transmembrane PRO polypeptide associated primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antisense oligonucleotide; cytostatic; immunosuppressive; antiinflammatory; gene therapy; hyperproliferative disorder; cancer; autoimmune; inflammatory disorder; inhibitor-kappa B kinase-gamma; ss;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New secreted and transmembrane nucleic acids and polypeptides, designated as PRO, useful for treating inflammation, organ failure, atherosclerosis, cardiac injury, infertility, birth defects, premature aging, AIDS, or
                                  06-OCT-2001; 2001US-00972607
                                                                                          03-OCT-2002; 2002WO-US031809
                                                                                                                                                                                                             WO2003031576-A2
                                                                                                                                                                                                                                                                                                                                                          modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 114; Page 187; 453pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                             /note= "
                                                                                                                                                                                                                    /*tag= c
/mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOB) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "Phosphorothioate linkages, all cytosines are
methylcytosine"
                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag=
                                                                                                                                                                                                                                                                                                                                                                                   /mod_base= OTHER
/note= "2'-methoxyethyl (2'-MOE) nucleotides"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        #ISIS 115419 #SEQ ID 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
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Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
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RESULT 907
AAL61645
ID AAL616
XX AAL616
XX AAL616
XX Iniol-
XX Nuclei
XX Nuclei
XX Unider
XX Unider
XX Unider
XX O9-OC:
PN 09-OC:
PR 07-DE(
XX OANO
PI Park (
XX Detec
PT havin
PT harnop
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to an antisense compound that is targeted to a cc nucleic acid encoding inhibitor-kappa B kinase-gamma, specifically chybridising to the nucleic acid encoding inhibitor-kappa B kinase-gamma cc and inhibiting its expression. Compounds of the invention are antisense coligonucleotides comprising at least one modified internucleoside coligonucleotides comprising at least one modified internucleoside coligonucleotides comprising at least one modified sugar moiety, which is a 2-0-methoxyethyl sugar moiety, or at least one modified nucleobase, which is a 5-methyleytosine. Preferably, the compound of cc thinternucleobase, which is a 5-methyleytosine. Preferably, the cc antisense oligonucleotide is a chimeric oligonucleotide. The compound of cc the invention is useful for preparing a composition for treating a cc hyperproliferative disorder e.g., cancer, or an autoimmune or cc inflammatory disorder. The methods are useful for inhibiting the cc expression of inhibitor-kappa B kinase-gamma in cells or tissues, and cc treating an animal having a disease or condition associated with cc inhibitor-kappa B kinase-gamma. Sequences given in ADA44713-ADA44790 cc inhibitor-kappa B kinase-gamma. Sequences given in ADA44713-ADA44790 cc inhibitor-kappa B kinase-gamma mRNA levels.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New compound having sequence targeted to nucleic acid encoding inhibitor-kappa B kinase-gamma, useful for preparing composition for treating e.g., cancer, or inflammatory or autoimmune disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAL61645 standard; DNA; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 BP; 4 A; 5 C; 7 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 3;
                                                                                                                                                                              09-OCT-2001; 2001US-0327864P.
07-DEC-2001; 2001US-00008978.
                                                                                                                                                                                                                                                                     01-MAY-2003
                                                                                                                                                                                                                                                                                                                                          Unidentified
                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid detection;
                                                                                                                                                                                                                                                                                                                                                                                                             Thiol-modified oligo #4 used in the nucleic acid detection method
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAL61645;
                                                                      WPI; 2003-430409/40.
                                                                                                                                           (NANO-)
                                                                                                                                                                                                                                  08-OCT-2002; 2002WO-US032088
                                                                                                                                                                                                                                                                                                         WO2003035829-A2
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                                                                                                                                             NANOSPHERE INC
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                                                                                                         Taton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                        TA,
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                                                                                                         Mirkin
                                                                                                                                                                                                                                                                                                                                                                               fabrication;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.2;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and

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allow hybridization,

and observing detectable

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RESULT 908
AL61472/c
ID AAL614
XX AAL614
XX Human
XX Human;
XW Human;
XW Human;
XW Human;
XW Iiver
XW WO2003
XW IISIS-NOV
XW 08-NOV
XW PA (ISIS-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a method of detecting a nucleic acid having two contions. The method involves providing nanoparticles having colligonuclectides attached to it which has a sequence complementary to sequence of two portions of nucleic acid, contacting nucleic acid and nanoparticles to allow hybridisation of oligonucleotides with two or more portions of nucleic acid and observing a detectable change brought about by hybridisation. The method and aggregate probes are useful for detecting two or more nucleic acids (from a biological source) having at least two portions such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic or structurally modified natural or synthetic RNA or DNA, or a product of applymerase chain reaction amplification. The invention is useful for preparing a nanoprobe conjugate for detecting an analyte and for detecting a nucleic acid bound to an electrode surface. It is also useful for fabrication and for separating a selected nucleic acid having two portions from other nucleic acids. The present sequence is an oligo used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 17
                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; activating transcription factor 3; ATP3; ischaemia; diabetes;
                                08-NOV-2001; 2001US-00010002.
                                                                 04-NOV-2002; 2002WO-US035331
                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                              modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL61472;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP;
                                                                                                                                  WO2003040161-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 18; Page 179;
   (ISIS-) ISIS PHARM INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATP3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                               /mod_base= OTHER
/note= "Phosphorothioate
methylcytidines"
                                                                                                                                      /*tag= c
/mod_base= OTHER
/mote= "2'methoxyethyl nucleotides"
                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                               /*tag= b
/mod_base= OTHER
/note= "2'methoxyethyl nucleotides"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   oligonucleotide,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0,
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Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LRF-1; antisense
kbone; antisense;
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                                                                                                                                                                                                                                                                                                                  backbone; All cytidines
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                therapy; CRG-5; LRG-21;
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to antisense compounds, compositions and methods for modulating the expression for activating transcription factor 3 (ATF3). ATF3 is also known as liver regeneration factor-1 (LRF-1), CRG-5, LRG-21, and TI-241. The invention is useful for the diagnosis, prevention and/or treatment of diseases or conditions associated with aberrant expression or activity of ATF3, such as ischaemia and diabetes. The entisense compound is useful in antisense therapy. The present sequence is an antisense oligonucleotide targetted to human ATF3 DNA. This sequence is used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                    Potato; plant; mitochondrial carrier protein; elongation factor transferrin binding protein; receptor-like protein kinase; helic non-long terminal repeat retroelement reverse transcriptase; overwatering; transgenic; reverse transcriptase; PCR; primer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antisense oligonucleotide compounds, useful for diagnosing, preventing and/or treating conditions with aberrant activity of activating transcription factor 3, such as ischemia and diabete
                                                                                                                                                                                                                      22-MAR-2001; 2001DE-01014063.
                                                                                                                                                                                                                                                22-MAR-2001; 2001DE-01014063
                                                                                                                                                                                                                                                                          10-OCT-2002
                                                                                                                                                                                                                                                                                                   DE10114063-A1
                                                                                                                                                                                                                                                                                                                                                                                                                     Potato gene
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ABZ59815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ABZ59815 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 20 BP; 7 A; 6 C; 4 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example 15; Page 77; 126pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Baker BF,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3958 ATGGTGGCAGGCCTCTGCT 3977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dobie K;
                                                                                                                                                                                                                                                                                                                                                                                                                     PCR primer dT20
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                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RNA;
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Pred. No. 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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and diabetes.
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The invention relates to DNA sequences (I) that encode six specific plant proteins: (i) a protein (ABP60425) with mitochondrial carrier protein activity (IIa), (ii) a protein (ABP60426) with transferrin binding protein activity (IIb); (iii) a protein (ABP60427) with receptor-like

New DNA sequences from potato, useful for producing plants with altered properties, e.g. tolerance of flooding, also related proteins, antibodis

Buelow L,

Techarntke

Į,

Haussuehl

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2003-041808/04.

1;

Page

German.

sequences. 8; 26pp;

(MPBC-) MPB COLOGNE GMBH MOLECULAR PLANT

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RESULT 910

ACA72109/c

ID ACA721

XX ACA721

XX ACA721

XX Human

XX Human;

KW Human;

KW erythr

KW apopto

KW jaftro

KW jaftro

KW jaftro

KW jamon

KW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CC protein kinase activity (IIc); (iv) a protein (ABB60428) with elongation CC factor EF-2 activity (IId); (v) a protein (ABB60429) with non-long CC terminal repeat retroelement reverse transcriptase activity (IIe); or CC (vi) a protein (ABB60430) with helicase activity (IIf). (I), also related CC sequences, derived ribozymes and antiense sequences, expression vectors, CC encoded proteins and antibodies against the proteins, are used to produce CC plants with altered properties, including tolerance of overwatering. The CC immunoassays. Also (I) or their primer or probe fragments and in CC inducible plant promoters and constitutively, aerobically or anaerobically CC inducible plant promoters, specifically for use in potatoes and the CC sequence that encodes (IId) is used to alter the translation profile in CC plants. Since (I) are derived from potato, their promoters and control endogenous genes. The present sequence is that of a PCR primer CC used in the first strand synthesis of cDNAs derived from Potato
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches 17;
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Best Local (
17-OCT-1997;
03-NOV-1997;
13-NOV-1997;
21-NOV-1997;
11-NAR-1998;
11-NAR-1998;
11-NAR-1998;
11-NAR-1998;
12-NAR-1998;
12-NAR-1998;
20-NAR-1998;
20-NAR-1998;
20-MAR-1998;
20-MAR-1998;
20-MAR-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          gastrointestinal disorder; Alzheimer's disease; Parkinson's disease; hypertension; myocardial ischaemia; kidney disease; carcinogenesis; glomerulonephritis; lung disease; pulmonary hypertension; preeclampsia; bronchial asthma; gastric ulcer; renal failure; cardiovascular disease; inflammatory bowel disease; reproductive disorder; premature labour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; ds; thrombolytic agent; interferon; interleukin; cytokine; erythropoietin; colony stimulating factor; cancer; colorectal carcinoma; apoptosis related condition; AIDS; amyotrophic lateral sclerosis; inflammatory disease; asthma; atherosclerosis; neurodegenerative disease;
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97US-0062250P
97US-0064249P
97US-0065311P
97US-0066364P
98US-0077632P
98US-0077641P
98US-0077641P
98US-0077641P
98US-0077044P
98US-0078004P
98US-007801P
98US-0078939P
98US-0078939P
98US-0078939P
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Pred. No. 9
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      26-MAR-1998
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30-NCV-1998
07-CCT-1998
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11-FEB-2000
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11-MAR-2000
                     2001US-00816744

2001US-00816920

2001WS-US009552

2001US-00854280

2001US-00872035

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2000WO-US008439.
2000WO-US013705.
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2000US-00709238.
2000US-00723749.
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2000WO-US023328.
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2000WO-US014941.
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2000WO-US005004.
2000WO-US005841.
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98US-0079664P.
98US-0079664P.
98US-0079786P.
98US-0079728P.
98US-0079728P.
98US-00168978.
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98US-00187368.
98US-00187368.
98US-00218517.
98US-00205445.
99US-00254465.
99US-00265186.
99US-00265186.
99US-00265186.
99US-00265186.
99US-00265180.
99US-00265180.
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99US-00380132.
99US-00380133.
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99US-0038855.
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RESULT 911
ABZ59177/c
ID ABZ591
XX ABZ591
XX ABZ591
XX ABZ591
XX CABZ591
XX G-prot
KW G-prot
KW TGR158
KW TGR158
KW Antico
KW Antico
XX Immuno
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            polypeptide, designated as PRO polypeptide. The PRO polypeptide is useful in PRO polypeptide detection methods. The PRO polypeptide is useful for linking a bioactive molecule to a cell. The PRO polypeptide or an antibody against it is useful for modulating a biological activity of a cell. The PRO polypeptide is useful in industrial applications including pharmaceuticals, diagnostics, biosensors and bioreactors. The PRO polypeptide is useful in industrial applications including polypeptide is also useful as a thrombolytic agent, interferon, interferon, interferon, as a colorectal carcinoma; apoptosis related conditions e.g. AIDS, cancer e.g. colorectal carcinoma; apoptosis related conditions e.g. AIDS, amyotrophic lateral sclerosis; inflammatory disease e.g. asthma, atherosclerosis; neurodegenerative disease e.g. Alzhaimer's disease, parkinson's disease; cardiovascular disease e.g. Alzhaimer's disease, cardiovascular disease e.g. Alzhaimer's disease, pastrointestinal disorders e.g. gastric ulcer and inflammatory bowel disease; reproductive disease e.g. renal failure and cardiovascular disease; reproductive disease e.g. pulmonary hypertension bronchial asthma; astrointestinal disorders e.g. gastric ulcer and inflammatory bowel disease; reproductive disease e.g. pulmonary hypertension. Note: The sequence data for this patent did not form part of the printed specification but was obtained in electronic format directly from USPTO at sequence data for this patent did not form part of the printed
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Best Local S
Matches 17
                                                                                                                                                                                                                                  G-protein coupled receptor; GPCR; TGRZ; TGRZO; .....TGR358; antiasthmatic; neuroprotective; cerebroprotective; nephrotenticonvulsant; hypotensive; hepatotropic; dermatological; human; anticonvulsant; hypotensive; hepatotropic; primer; ss.
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Goddard
Kljavin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New isolated PRO polypeptides e.g. PRO213, PRO274 and PRO300, for use as pharmaceuticals, diagnostics, biosensors and bioreactors, for identifying modulators of receptor-ligand interactions.
                                                                    16-JAN-2003.
                                                                                                                                                                                                                                                                                                                                                                                                                Human TGR118 DNA expression analysing RT-PCR forward primer.
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                                                                                                                                WO2003004678-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   invention relates to an isolated secreted and transmembrane
ypeptide, designated as PRO polypeptide. The PRO polypeptide is useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5196 TCAGCGTGGGAGGCCACGTG 5215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2003-328499/31.
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17; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         J, Baker KP, Botstein D, Desnoyers L, Filvaroff E, Fong S, Gao W, Gerber H, Godowski FJ, Grimaldi JC, Gurney AL, I Kuo SS, Napier MA, Pan J, Paoni NF, Tumas D, Williams PM, Wood WI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4 A;
                                                                                                                                                                                                                                                                                                                                                  receptor; GPCR; TGR2; TGR38; TGR118; TGR164; TGR343;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7 C; 4 G; 5 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      577; 55pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <u>.</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 15.2;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1; Length 20
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I, Gerritsen ME;
Hillan KJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Roy MA,
                                                                                                                                                                                                                                                                                                                       nephrotropic;
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01-JUL-2002; 2002WO-US020860.

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ARSSULT 912
ARSS2749/C
ID ABX9277
XX
ARS 2749/C
ID ABX927
XX
DE Human;
XW Human;
XW Human;
XW immune
XW cardia
XW genetiar
XW cardia
XW genetiar
XW cardia
XW Jenetiar
XW US2002
XX 16-0CT
XX
PP 11-0CT
PR 13-NOV
PR 13-NOV
PR 13-NOV
PR 13-NOV
PR 13-NOV
PR 13-NOV
PR 21-NOV
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03-NOV-1997;
13-NOV-1997;
21-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention provides new G-protein coupled receptor (GPCR) polypeptides designated TGR2, TGR38, TGR118, TGR164, TGR343 and TGR358 and encoding polymelectides. The polypeptides can be expressed by standard DNA recombination methodology. The polypeptides are useful for screening or identifying modulators of GPCR or signal transduction. The modulators of signal transduction are useful for treating or preventing TGR-associated disorders, e.g. asthma, multiple sclerosis or kidney disease. The polypeptides are useful as targets for diagnosing or treating e.g. epilepsy, stroke, neurodegeneration, hypogonadism, hyperprolactinemia, asymptomatic urinary abnormalities, hypertension, nephrolithiasis, cirrhosis, lesions, jaundice, psooriasis, lupus erythematosus, or acute inflammatory dermatoses. Sequences ABZ59173-184 represent RT-PCR primers used for analysing the expression of the various TGR polypeptides in
                                                                                                                                                                                                                                            genetic disorder; cytostatic; antidiabetic; antiinflammatory; antiatthritic; anti-tumour; vulnerary.
                                                                                                                                                                                                                                                                                                  Human; PRO polypeptide; secreted and transmembrane protein; immune disorder; diabetes; hyper-insulinaemia; hypo-insulinaemia; cardiac insufficiency; nervous system disorder; kidney disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New G-protein coupled receptor polypeptides designated TGR2, TGR38, TGR118, TGR164, TGR343 and TGR358, useful as targets for screening for treating or preventing e.g. asthma, multiple sclerosis, stroke
                                                   26-MAY-1981;
17-OCT-1997;
                                                                                                                                    14-NOV-2002.
                                                                                                                                                                                                                                                                                                                                                                   Human PRO DNA PCR primer SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                      20-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABX92749 standard; DNA; 20
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                                                                                                    16-OCT-2001;
                                                                                                                                                                    US2002169284-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1431 TGTGGAGGAGAATCGAGGAC 1450
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                    2001US-00978697.
 97US-0064249P.
97US-0065311P.
97US-0066364P.
                                                   97US-0062250P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВÞ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 15.2; DB 1
Pred. No. 9.3e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 20;
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110-MAR-1998
111-MAR-1998
111-MAR-1998
111-MAR-1998
113-MAR-1998
120-MAR-1998
20-MAR-1998
20-NOV-1998
20-NOV-1998
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10-MAR-1999
10-MAR-1999
10-MAR-1999
10-MAR-2000
10-MAR-2000
11-FEB-2000
21-MAR-2000
21-MAR-2000
21-MAR-2000
21-MAR-2000
22-MAR-2000
21-MAR-2000
98US-0077450P
98US-0077643P
98US-0077643P
98US-0077649P
98US-0077649P
98US-007794P
98US-0078930P
98US-0078930P
98US-0079663P
98US-0079663P
98US-0079663P
98US-0079663P
98US-0079663P
98US-0079663P
98US-0079663P
98US-00105413
98US-00105413
98US-00105413
98US-00105413
98US-00184216
98US-00184216
98US-00202654
98US-0031183
98US-00202654
99US-0030376
99US-0030376
2000WO-US003274
2000WO-US003276
20
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RESULT 913
ABX79181
ID ABX7918
XX
AC ABX791
XX
Thio-n
XX
DE Thio-n
XX
Nanopa
KW harpan
KW herpse
KW Epstei
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                                                                                                                                                                                                                                                                                                                                                                                           밁
                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention relates to the isolation of novel human PRO CC polypeptides, and the polynuclectide sequences encoding them. The PRO CC polypeptides are useful for detecting other PRO polypeptides, for linking CC bioactive molecules to cells expressing PRO polypeptides, for modulating CC bioactive molecules to cells expressing PRO polypeptides, for modulating CC biological activities of cells expressing PRO polypeptides, and for for CC identifying agonists or antagonists. The bioactive molecule maybe a CC toxin, radiolabel or antibody, and causes apoptosis or death of the cell. The PRO polypeptides are useful for treating immune disorders, diabetes CC or hyper- or hypo-insulinaemia, cardiac insufficiency, nervous system CC tumours, and wound healing. The polypucleotide sequences encoding PRO CC tumours, and wound healing. The polypucleotide sequences encoding PRO CC polypeptides are useful as hybridisation probes, in chromosome and gene CC mapping, in the generation of antisense RNA and DNA, in the preparation CC animals, for the genetic analysis of individuals with genetic disorders, CC and in gene therapy. The present sequence represents a PCR primer used in the examples of the present invention. Note: The sequence data for this car are septiment was obtained in electronic format directly from the USPTO web site
                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 17
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Ferrara N, I
Goddard A, (
Kljavin IJ,
Stewart TA,
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01-JUN-2001;
01-JUN-2001;
05-JUN-2001;
14-JUN-2001;
19-JUN-2001;
20-JUN-2001;
29-JUN-2001;
09-JUN-2001;
30-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel secreted and transmembrane polypeptides and encoding them useful for treating cancer, kidney cartilage disorders and immune deficiencies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI;
                                                                                                                                                                                                                                                                                                                                                                                           Sequence 20
                                                                                                                                                                               ABX79181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 114; Page 192; 459pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GETH )
                                                                                                                                                                                                                                                                                                                                                                                                                         seqdata.uspto.gov/psipsDIDEntry.html
                                                                                                                                                                                                                                                                               5196 TCAGCGTGGGAGGCCACGTG 5215
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                                                                                                                                                                                                                                                                                                                         17;
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                                                                                                                                                                                                                                                                                                                                            Similarity
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                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                              TCAGTGTGAAAGGCCACGTG 1
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Filvaroff E,
Godowski PJ, (
Kuo SS, Napi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001WO-US017092.
2001WS-00872035.
2001WS-00874503.
2001US-00874503.
2001US-00882636.
2001WS-00886342.
2001WS-00886342.
2001WS-00886342.
2001WS-US021066.
2001WS-US021066.
                                                                                                                                                                                                                                                                                                                                                                                           BP; 4 A; 7 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                          0.3%;
                                                                                                                                                                               DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KP, Botstein D, Desnoyers L, Eaton D;
f E, Fong S, Gao W, Gerber H, Gerrisen ME;
PJ, Grimaldi JC, Gurney AL, Hillan KJ;
Napler MA, Pan J, Paoni NF, Roy MA, Shelt,
Williams PM, Wood WI;
                                                                                                                                                                               20
                                                                                                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                            Score 15.2; DB 1
Pred. No. 9.3e+02
                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                             Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     diseases,
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diseases, bone,
                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                              20;
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                                                                                                                                                                                                                                                                                                                              Gaps
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Nanoparticle; ss; nucleic acid detection; viral disease; probe; human immunodeficiency virus infection; hepatitis virus infection; herpes virus infection; cytomegalovirus infection; forensic scienc Epstein-Barr virus infection; bacterial disease; gene therapy;

science;

Thio-modified 15-APR-2003

20dA oligonucleotide.

(first

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RESULT 914
ACC47072/c
ID ACC47072 standard; DNA; 20
XX
AC ACC47072;
XX
DT 05-JUN-2003 (first entry)
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                                                                                                                                  Matches
                                                                                                  Query Match
Best Local
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21-JUL-1997;
29-JAN-1999;
                                                                                                                                                                                                                                                                                                                          Detecting nucleic acids having at least 2 portions comprises use nanoparticles which have oligonucleotides attached to them that complementary to portions of the nucleic acid sequence.
                                                                                                                                                                                                                                                                                                                                                                                                        29-JAN-1999; 99US-00240755.
25-JUN-1999; 99US-00344667.
26-APR-2000; 2000US-0200161P.
26-JUN-2000; 2000US-00603830.
                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                             Example 18; Page 44; 130pp; English.
                                                                                                                                                                                                                                                                                                                                                                            Mirkin CA,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          12-OCT-2001; 2001US-00976577.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sexually transmitted disease; inherited disorder; DNA sequencing;
paternity testing; cell line authentication.
                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-198491/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US2002155462-A1
                                                                                                                                                                                                                                                                                                                                                                                          (NANO-) NANOSPHERE INC.
                                                                             5393 AAAAAAATACAAAAAAGAAA 5412
                                                                                            17; Conserv
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                                                                AAAAAAAAAAAAAAAAA 20
                                                                                                                       20
                                                                                                                       BP;
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                                                                                            Conservative
 (first entry)
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97WO-US012783.
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Pred. No. 9.3e+02
                                                                                                                       G; 0 T; 0 U; 0 Other;
                                                                                            Mismatches
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                                                                                                        DB 1; Length
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TGCCTGCCTGCTTCCTTCAC

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The present invention describes a compound (I) comprising 8-50 cnucleobases which is targeted to a 5 untranslated region (UTR), coding, CC introduced which is targeted to a 5. Untranslated region (UTR), coding, CC if the compound specifically hybridises with care introduced phospholipase and inhibits the expression of phospholipase A2, group IIA (synovial). CC Also described: (1) a composition comprising the compound and a carrier CC or diluent; (2) a method of inhibiting the expression of phospholipase CC A2, group IIA in cells or tissues; and (3) a method of treating an animal compound (I) has antiinflammatory, antidiabetic, cytostatic and CC (synovial). (I) has antiinflammatory, antidiabetic, cytostatic and CC antipsoriatic activities, and can be used in vaccines and in gene compound (I) can be used for preparing a composition for treating or preventing inflammation, cancer, psoriasis or diabetes. The present sequence represents a mouse phospholipase A2 group IIA (synovial) content of the compound (I) can be used for preparing a composition for composition for the compound (I) can be used for preparing a composition for composition for the compound (I) can be used for preparing a composition for composition for the compound (I) can be used for preparing a composition for composition for the compound (I) can be used for preparing a composition for composition for the compound (I) can be used for preparing a composition for composition for the compound (I) can be used for preparing a composition for composition for the compound (I) can be used for preparing a composition for composition for the compound (I) can be used for preparing a composition for composition for the compound (I) can be used for preparing a composition for composition for the compositi
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New compound that hybridizes with and inhibits the expression of phospholipase A2, group IIA, useful for preparing a composition treating or preventing inflammation, cancer, psoriasis or diabet
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Phospholipase A2 group IIA; synovial; antisense modulation; inflammatio phospholipase A2 group IIA inhibitor; phosphorothioate; antiinflammator antidiabetic; cytostatic; antipsoriatic; vaccine; gene therapy; cancer;
                                                                                                                                                                                                                                                                                                 example from
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                                                                                                            Local
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                                                                        l Similarity
17; Conserv
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TGCCTGCCTGATACCCTCAC 477
                                                                                                                                                                                                                         BP; 7
                                                                        Conservative
                                                                                                                                                                                                                                                                                                     the present invention
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16. .20
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/mod_base= OTHER
/note= "2'-O-methoxyethyl (2'-MOB) gapmer"
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/mod_base= OTHER
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/mod_base= OTHER
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                                                                                                                Score 15.2;
Pred. No. 9
                                                                            Mismatches
                                                                                                                9.3e+02
                                                                                                                                                    DB 1;
                                                                                                                                                Length 20;
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                                                                            Gaps
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RESULT 915
ACA66490;
ACA66490;
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ACA66490;

17-OCT-1997
13-NOV-1997
13-NOV-1997
11-NOV-1997
11-NAR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15-OCT-2001;
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97US-0062250P

97US-006314P

97US-0063364P

98US-0077641P

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98US-0077641P

98US-00778910P

98US-0078910P

98US-0078910P

98US-0078936P

98US-0079664P

98US-00797294P

98US-0079728P

98US-0079728P

98US-0079728P

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98US-00168978

99US-0031813

99US-00380138

99US-00380138

99US-00388551

99US-00388551
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16-DEC-1999
30-DEC-1999
30-DEC-1999
30-DEC-1999
06-JAN-2000
06-JAN-2000
06-JAN-2000
06-JAN-2000
01-MAR-2000
01-MAR-2000
01-MAR-2000
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99WO-US031243.
99WO-US031274.
2000WO-US000219.
2000WO-US0003565.
2000WO-US005841.
2000WO-US005841.
2000WO-US005841.
2000WO-US0065841.
2000WO-US0065841.
2000WO-US0065841.
2000WO-US0065841.
2000WO-US014942.
2000WO-US014942.
2000WO-US014941.
2000WO-US030871.
2001WS-00816740.
2001WS-00854280.
2001WS-00854280.
2001WS-0089263.
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2001WS-0089263.
2001WS-0089263.
2001WS-00886342.
2001WS-0US017969.
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2001WS-US017969.
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2001WS-US017969.
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(GETH _ GENENTECH INC.

Goddard Kljavin Stewart Ferrara Ashkenazi i i i Ŗ, Godowski PJ, 「, Kuo SS, Nar , Tumas D, Wi Filvaroff E, Napier MA, Pa Williams PM, Botstein D, Desnoyers L, Eaton DL; Fong S, Gao W, Gerber H, Gerriten Grimaldi JC, Gurney AL, Hillan KJ; ier MA, Pan J, Paoni NF, Roy MA, Sh lliams PM, Wood WI; Shelton DL; ă

2003-341189/32.

New genes and secreted and tra PRO1559), useful for treating atherosclerosis, infertility, sclerosis in mammals. transmembrane polypeptides (e.g. PRO337 ing or diagnosing e.g. cancers, ty, stroke, encephalitis, hepatitis or m or multiple õ

Example 114; Page 193; 460pp; English.

The invention relates to a new isolated nucleic acid molecule comprises a sequence with at least 80% identity to: (a) a nucleotide encoding any of 94 PRO polypeptides whose sequences are fully defined in the specification; or (b) any of 94 nucleotide sequences fully defined in the specification; or the full length coding sequence of any these 94 nucleotide sequences. Also included are an isolated PRO polypeptide scoring at least 80% positives when compared to any of the PRO polypeptide sequences cited above (or an isolated PRO polypeptide having at least 80% amino acid sequence identity to: (a) an amino acid sequence encoded by the nucleotide deposited with ATCC numbers listed in the specification; (b) the PRO polypeptide, lacking its associated signal peptide; or (c) an extracellular domain of the PRO polypeptide, with or lacking its associated signal peptide).

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RESULT 916
ABX92177
ID ABX921
XX ABX921
XX Nanop
XX Nanop
XX Nonop
XX Nonop
XX Nonop
XX Nonop
XX Hepat
XX bact
XX bact
XX bact
XX Synt
XX 29-
PN 29-
PR 29-

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local (
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Taton TA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nonoparticle; nucleic acid detection; hybridisation; diagnosis; sequencing; viral infection; human immunodeficiency virus; HIV; hepatitis virus; hepatitis virus; hepatitis lifection; sexually transmitted disease; inherited disorder; forensic; paternity testing; cell line authentication; gene therapy; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-JUL-1996;
21-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        (NANO-) NANOSPHERE INC.
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97WO-US012783.
99US-00240755.
99US-00344667.
2000US-0200161P.
2000US-00603830.
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                                                                                                                                                                                                                                                                                                                                                             Storhoff JJ,
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Detecting nucleic acids having at least two portions involves use of nanoparticles which have oligonucleotides attached to them that are complementary to portions of the nucleic acid sequence.

14-MAY-2001;

(UYQU-) UNIV

QUEENSLAND TECHNOLOGY 2001AU-00004971. 14-MAY-2002; 2002WO-AU000591

21-NOV-2002

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           밁
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Best Local
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                                                                                                                                                                                                                                                                                                                                                      WO200292629-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
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diagnostic marker; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-MAY-2003
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17; Conserv
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Pred. No. 9.
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RESULT 918
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Matches 17
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                     The invention relates to a novel method for detecting a point mutation in a DNA strand. The novel method comprises hybridising a target DNA strand containing or suspected of containing a point mutation with a test nucleic acid strand complementary to the DNA strand to form a target DNA strand/test nucleic acid strand duplex, contacting the duplex with an strand/test nucleic acid strand duplex, contacting the duplex with an RNase H, and determining whether the ribonucleotide residues within the
                                                                                                                                                                                                                                                                                duplex
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAR-2001; 2001US-00823634
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                                                                                                                                                                                                                                                                                ing point mutation in DNA strand, mutation with test DNA strand to with RNase H and determining the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleotide
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TSENG T.
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                                                                                                                                                                                                        Fig
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form duplex, contacting the
cleavage of test strand by RNase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        target
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Matches 17; Conserv
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21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
26-APR-2000;
26-JUN-2000;
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                                                                                                                                             comprises
them that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      virus detection; human immunodeficiency virus; HIV; hepatitis; herpes; cytomegalovirus; Epstein-Barr virus; bacterial disease; DNA sequencing sexually transmitted disease; inherited disorder; forensic; paternity testing; cell line authentication.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACD27255 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                               Example 18;
                                                                                                                                                                                          Detecting
                                                                                                                                                                                                                                                                                            Taton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      24-OCT-2002.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nanotechnology; ss; nucleic acid detection; nanoparticle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nanotechnology nucleic acid detection method associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-OCT-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACD27255;
                                                                                                                                                                                                                                                                                                                                                                (NANO-) NANOSPHERE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5393
                                                                                                                                                                                                                                            2003-228114/22.
                                                                                                                                                                                                                                                                                            TA,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAAAAATATTAAAAAAAA 20
                                                                                                                                          nucleic acids having 2 portions e.g. for detecting disease, use of nanoparticles which have oligonucleotides attached to are complementary to portions of the nucleic acid sequence.
                                                                                                                                                                                                                                                                                                                                                                                                              ; 96US-0031809P.
; 97WO-US012783.
; 99US-00240755.
; 99US-00344667.
; 2000US-0200161P.
; 2000US-00603830.
                                                                                             Page 43; 129pp;
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                                                                                                                                                                                                                                                                                                                   Letsinger
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /*tag=
/mod_ba
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "OTHER= Thiol modified"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; 0 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.3%;
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                                                                                                                                                                                                                                                                                                                   RĽ,
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                                                                                                                                                                                                                                                                                                                     Mucic
                                                                                             English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 15.2;
Pred. No. 9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                   RC,
                                                                                                                                                                                                                                                                                                                     Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1;
).3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                   Elghanian R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequencing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ç
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This invention relates to a novel method for detecting a nucleic having 2 portions. The method comprises providing nanoparticles beligonucleotides attached, where the oligonucleotide on each nanoparticles.

les having nanoparticle

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RESULT 920
ACD27125
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Best Local (
                                                                                                                                                                                                                                                                     29-JUL-1996;
21-JUL-1997;
29-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nanotechnology
                                                                             Mirkin CA,
                                                                                                                                                                                                                   25-JUN-1999;
26-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US2002164605-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nanotechnology; nucleic acid detection; nanoparticle; 88; forensic; DNA sequencing; paternity testing; cell line authentication.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20 BP; 20 A; 0 C; 0
                                                          Taton
                                                                                                                                                                                                                                                                                                                                                                               28-SEP-2001; 2001US-00966312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_base
WPI; 2003-247253/24
                                                                                                                                        (NANO-) NANOSPHERE INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAAAAAAAAAAAAAAAA 20
                                                                                                                                                                                    96US-0031809P.

97WO-US012783.

99US-00240755.

99US-00344667.

2000US-0200161P.

2000US-00603830.
                                                                                   Letsinger
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid detection method oligonucleotide #54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mod_base= OTHER
note= "OTHER= Thiol modified" "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             *tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.3%;
                                                                                   RL,
                                                                                   Mucic RC,
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Pred. No. 9.3e+02;
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                                                                                   Storhoff JJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Length 20;
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                                                                                   Blghanian
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FFFFX8X5555555555555555555555555555555
                                                                                                                                                                                                    Detecting nucleic acid having two portions, by providing nanoparticles having oligonucleotides attached to it, contacting nucleic acid and nanoparticles to allow hybridization, and observing detectable change, useful in forensics.
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Example 18; Page 44; 130pp; English.

Requences having two portions. The method involves providing contacticles having oligonucleotides attached to them, which has a contacting nucleic acid and nanoparticles, to allow hybridisation of contacting nucleic acid and nanoparticles, to allow hybridisation of contacting nucleic acid and nanoparticles, to allow hybridisation of contacting nucleic acid and nanoparticles, to allow hybridisation of contacting nucleic acid and nanoparticles, to allow hybridisation of convention and the aggregate probes are useful for detecting two or more nucleic acids (from a biological source) having at least two portions, such as viral RNA or DNA, bacterial or fungal DNA, a gene associated with a disease, synthetic, or structurally-modified natural or synthetic RNA or DNA, or a product of a polymerase chain reaction amplification.

CC NANOPARTICLES and nanoparticle-oligonucleotide conjugates of the nucleic acid having two portions from other nucleic acids. The method of the invention is useful in forensics, DNA sequencing, for paternity capting, cell line authentication, and monitoring gene therapy. Diagnostic assays employing the nanoparticle-oligonuclectide conjugates of the invention improve the sensitivity of the nucleic acid detection assay. The present sequence represents a thiol modified oligonuclectide sequence used to demonstrate the method of the invention This invention relates to a novel method for detecting nucleic acid

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                                           Query Match
Best Local S
Matches 17
                                                                                      Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
             5393 AAAAAAAATACAAAAAAAGAAA 5412
|||||||| | |||||| |||
                                           17; Conserv
AAAAAAAAAAAAAAAAA 20
                                           0.3%;
ilarity 85.0%;
Conservative
                                           0; Mismatches
                                                      Score 15.2;
Pred. No. 9.
                                                      DB 1;
                                                                  Length
                                              Indels
                                                                   20
                                              0
                                              Gaps
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RESULT 921
ACD27385
ID ACD273
XX ACD273
XX ACD273
XX ACD273
XX Nanote
XX Nan
29-JUL-1996;
21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
26-APR-2000;
26-JUN-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        pathogen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nanoparticle; ss; nucleic acid detection; DNA sequencing;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nanotechnology nucleic acid detection method associated #54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ACD27385
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ACD27385 standard; DNA; 20
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    99US-00344667.
2000US-0200161P.
2000US-00603830.
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                                                                                                                           96US-0031809P.
97WO-US012783.
99US-00240755.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /*tag= a
/mod_base= OTHER
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RESULT 922
ACD27190
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention relates to a novel method for detecting a nucleic acid by CC contacting a nucleic acid with at least two types of nanoparticles having CC oligonucleotides attached, allowing hybridisation of the oligonucleotides CC on the nanoparticles, and observing a detectable change. The CC oligonucleotides on each nanoparticle have a sequence complementary to CC its respective portion of the sequence of the nucleic acid to be CC detected. The method of the invention may be used for the detection of a CC nucleic acid used in, e.g. research and analytical laboratories in DNA CC sequencing, in the field to detect the presence of specific pathogens, in CC the doctor's office for quick identification of an infection to assist in CC prescribing a drug for treatment, and in homes and health centres for CC inexpensive first-line screening. The method of the invention detects nucleic acids based on observing a colour change with the naked eye. This CC method is cheap, fast, simple, robust and does not require specialised or cxpensive equipment. The present sequence represents a thiol modified coligonucleotide sequence used to demonstrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 17
29-JUL-1996;
21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
26-APR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detection of nucleic acid for, e.g. research in deoxyribonucleic acid sequencing, involves with nanoparticles having oligonucleotides.
                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                            Nanoparticle;
                                                                                                                                                                                                                                                                                                                                         Nanotechnology nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                        ACD27190
                                                                                                                                                                                                                                                                                                                                                                                                                                   ACD27190 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mirkin CA,
                                                                                            12-OCT-2001;
                                                                                                                                                        US2002182613-A1
                                                                                                                                                                                                                                modified_base
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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96US-0031809P.
97WO-US012783.
99US-00240755.
99US-00344667.
2000US-0200161P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                           2001US-00976971.
                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                            ss; nucleic acid detection; DNA sequencing
                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                    /*tag=
                                                                                                                                                                                      note=
                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.3%;
                                                                                                                                                                                                      base= OTHER
                                                                                                                                                                                      "OTHER= Thiol modified"
                                                                                                                                                                                                                                                                                                                                                                                                                                    20
                                                                                                                                                                                                                                                                                                                                                                                                                                    ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                           detection method associated #54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 15.2; DB 1; Length Pred. No. 9.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and analytical laboratories contacting nucleic acid
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SXCCCCCCCCCCCCCCCXSXFFFXXXXFFFXXXXXFF
                                                         contacting nucleic acid with at least two types of nanoparticles having oligonucleotides, allowing hybridisation of the oligonucleotides on the nanoparticles, and observing a detectable change. The oligonucleotides on the cach nanoparticle have a sequence complementary to its respective portion of the sequence of the nucleic acid. The method of the invention may be used for the detection of a nucleic acid used in, e.g. research and analytical laboratories in DNA sequencing, in the field to detect the presence of specific pathogens, in the doctor's office for quick identification of an infection to assist in prescribing a drug for iteratment, and in homes and health centres for inexpensive first-line screening. The inventive method of detecting nucleic acids based on observing a colour change with the naked eye are cheap, fast, simple, robust (the reagents are stable), do not require specialised or expensive equipment, and little or no instrumentation is required. The present sequence represents a thiol modified oligonucleotide sequence used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Detection of nucleic acid for, e.g. research and analytical laboratories in deoxyribonucleic acid sequencing, involves contacting nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-596265/56.
Sequence 20
                                              demonstrate
                                                                                                                                                                                                                                                                                                                                                                                                         This invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mirkin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NANO-) NANOSPHERE INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          26-JUN-2000; 2000US-00603830.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nanoparticles having oligonucleotides.
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  BP; 20
                                              the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Letsinger RL,
                                                                                                                                                                                                                                                                                                                                                                                                         relates
  A; 0 C; 0
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                                                                                                                                                                                                                                                                                                                                                                                                      a novel method for detecting a nucleic acid by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mucic RC,
  G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Storhoff JJ,
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Ś Query Match Best Local S Matches 17 5393 ААААААТАСАААААСААА 5412 Similarity AAAAAAAAAAAAAAAAAA 20 Conservative 0.3%; 0; Score 15.2; Pred. No. 9. Mismatches .3e+02; 멂 1; Length 20; Indels 0 Gaps 0

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                                                                                                                                                                                                                           RESULT 923
29-JUL-1996;
21-JUL-1997;
29-JAN-1999;
                                                                                                                                                          Nanotechnology nucleic acid detection method oligonucleotide #54.
                                                                                                                                           Nanotechnology; nucleic acid detection; nanofabrication; nanoprobe; ss
                                               06-MAR-2003
                                                               US2003044805-A1
                                                                                                    modified_base
                                                                                                                                                                            15-OCT-2003
                                                                                                                                                                                           ACD27060
                                                                                                                                                                                                           ACD27060 standard;
                               15-OCT-2001;
                               2001US-00981344
                                                                                                                                                                            (first
96US-0031809P.
97WO-US012783.
99US-00240755.
                                                                                    /*tag= a
/mod_base=
                                                                                                             Location/Qualifiers
                                                                             'mod_base= OTHER
'note= "OTHER= T
                                                                                                                                                                                                           DNA;
                                                                                                                                                                            entry)
                                                                               "OTHER= Thiol modified"
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RESULT 924
ACHOOG4
ID ACHOOG
XX ACHOOG
XX ACHOOG
XX ACHOOG
XX ACHOOG
XX Nanote
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          method comprises providing nanoparticles with oligonuclectides attached to them, which have a sequence complementary to a sequence of two comprises providing acid, contacting the nucleic acid and nanoparticles contacting the oligonucleotides with two or more portions contacting the present the set that when the change brought about by the hybridisation. The nucleic acid to be detected must have at least two portions and the distances between these are chosen so that when the canoparticle-oligonucleotide conjugate binds the target sequence a detecting two or more nucleic acids (from a biological source) having at cleast two portions, such as viral RNA, bacterial or fungal DNA, a gene associated with a disease, synthetic, or structurally-modified natural conspicies and nanoparticle-oligonucleotide conjugates of the invention and conspicies and proparticle conjugate for detecting an analyte, and for detecting a nanoprobe conjugate for detecting an analyte, and for detecting a nucleic acid bound to an electrode surface.

Conspicutions from other nucleic acids Diagnostic assays employing two particles acid detection methods and can be used to detect nucleic acids that are present in only small amounts in a sample. The invention also provides highly desirable nanoparticle-oligonucleotide conjugates. These conjugates are stable with tailoranticleotide sequence used to
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Best Local S
Matches 17
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Taton TA;
                                                                                                                                                       Nanotechnology nucleic acid detection method oligonucleotide #54.
                                                                                                                                                                                                   15-OCT-2003
                                                                                                                                                                                                                                          ACH00064;
                                                                                                                                                                                                                                                                               ACH00064 standard; DNA; 20 BP
            Key
modified_base
                                                                              Synthetic
                                                                                                                  Nanotechnology;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 18; Page 44; 130pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detection of nucleic acid having -2 portions used to prepare biomaterials and in nanofabrication methods, comprises providing nanoparticles,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-APR-2000;
26-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence represents a thiol modified oligonucleotide sequence used demonstrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2003-521746/49
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                 AAAAAAAAAAAAAAAAA 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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2000US-0200161P.
2000US-00603830.
                                                                                                                  nucleic acid detection; nanofabrication; nanoprobe; ss
                                     Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   slates to a novel method for detecting nucleic providing nanoparticles with oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 15.2;
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C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 20;
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ACH03110/ ID ACHO XX

ACH03110 standard; DNA; 20

ВP

RESULT 925

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5393 AAAAAAATACAAAAAAGAAA 5412

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CC to allow hybridisation of the oligonucleotides with two or more portions of the nucleic acid, and observing a detectable change brought about by CC the hybridisation. The nucleic acid to be detected must have at least two CC portions and the distances between these are chosen so that when the CC nanoparticle-oligonucleotide conjugate binds the target sequence a CC detecting two or more nucleic acids (from a biological source) having at CC least two portions, such as viral RNA, bacterial or fungal DNA, a gene CC associated with a disease, synthetic, or structurally-modified natural CC or synthetic RNA or DNA, or a product of a polymerase chain reaction CC amplification. Nanoparticle-oligonucleotide conjugates of the invention CC and for detecting a nucleic acid bound to an electrode surface.

CC Nanoparticles and nanoparticle conjugates of the invention are useful for separating a selected mucleic acid having two CC portions from other nucleic acids bound to an electrode surface.

CC nanoparticle-oligonucleotide conjugates improve the sensitivity of nucleic acid detection methods and can be used to detect nucleic acids to conjugates are present in only small amounts in a sample. The invention also provides highly desirable nanoparticle-oligonucleotide conjugates. The present conjugates are stable with tailored hybridisation abilities. These conjugates are stable with tailored hybridisation abilities. The present conjugates at this modified cliconucleotide sequence used to
  Matches
                    Query Match
Best Local (
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21-JUL-1997;
29-JAN-1999;
25-JUN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention relates to a novel method for detecting nucleic acids. The method comprises providing nanoparticles with oligonucleotides attached to them, which have a sequence complementary to a sequence of two portions of nucleic acid, contacting the nucleic acid and nanoparticles
                                                                             Sequence 20 BP; 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 18; Page 44; 108pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Detection of nucleic acid having at least two portions, by contacting nucleic acid and nanoparticles under conditions, which allows hybridization of oligonucleotides on nanoparticles with at least two
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2003-634854/60
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26-JUN-2000;
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                                                                                                                                  sequence represents a thiol modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                portions of nucleic acid.
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                    Similarity
                                                                                                                    the method of the invention
    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000US-0200161P
2000US-00603830
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97WO-US012783.
99US-00240755.
99US-00344667.
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/note= "OTHER= Thiol modified"
                                                                             A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                  85.0%;
                                          0.3%;
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0;
                    Score 15.2;
Pred. No. 9.
    Mismatches
                                                                                                                                    oligonucleotide sequence
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                      9.3e+02
                                      DB 1; Length 20;
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RESULT 926
ACD99851
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XX ACD998
AC ACD998
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XX Z5-SBI
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Treating non-allergic inflammatory diseases, such as psoriasis, ecz allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 BP; 2 A; 8 C; 3 G; 7 T; 0 U;
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                  US2003050268-A1
                                                                             Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antilicer; gene therapy; vaccine; non-allergic inflammatory disease; psoriasis; eczema; allergic contact dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
                                                                                                                                                                                                                                                        ACD99851 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 29; 229pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-521815/49
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17; Conserv
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                                                                                                                                                                                                                                                                                                                                       AGGATCAGGAGCGACATGGA 1
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Pred. No. 9.3e+02
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Best Local S
Matches 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Treating non-allergic inflammatory diseases, such as psoriasis, ecz allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 20 BP; 20 A; 0 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAR-2002; 2002US-00112653
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                                                                                                                                                                                                                                                                                          Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergic inflammatory disease psoriasis; eczema; allergic contact dermatitis; latex dermatitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 23; 229pp; English.
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                                         WPI; 2003-521815/49.
                                                                  Krieg AM,
                                                                                                                                       29-MAR-2001; 2001US-0279642P
                                                                                                                                                                  29-MAR-2002; 2002US-00112653.
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                                                                                                                                                                                                                                                                              inflammatory bowel
                                                                                                                                                                                                                                                                                                                                               Immunostimulatory nucleic acid #533
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                                                                                                                                                                                                                                                                                                                                                                                                                                  ACD99847 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
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                                                                                             (KRIE/) KRIEG A M. (BERG/) BERG D J.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       <u>АААААААААААААА</u> 20
                                                                  Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Berg DJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                             therapy; vaccine; non-allergic inflammatory disema; allergic contact dermatitis; latex dermatitis allergic contact dermatitis; latex dermatitis wel disease; ulcerative colitis; Crohn's disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.3%;
                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15.2;
Pred. No. 9.
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Treating non-allergic inflammatory allergic contact dermatitis, latex

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                                                                       The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory disease, such as pooriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention describes a method of treating non-allergic inflammatory disease comprising administering to a subject having or at risk of developing a non-allergic inflammatory disease an immunostimulatory nucleic acid for prevention or treatment of the disease. The method is useful for treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact demmaticis, latex dermatitis or inflammatory bowel disease e.g., ulcerative colitis or Crohn's disease. This sequence represents an immunostimulatory nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Treating non-allergic inflammatory diseases, such as psoriasis, eczema, allergic contact dermatitis, latex dermatitis or inflammatory bowel disease by administering an immunostimulatory nucleic acid.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      antiulcer; gene the psoriasis; eczema; inflammatory bowel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACD99532 standard; DNA; 20 BP
                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAR-2001; 2001US-0279642P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAR-2002; 2002US-00112653
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13-MAR-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Immunostimulatory; antiinflammatory; dermatological; antipsoriatic; antiulcer; gene therapy; vaccine; non-allergot inflammatory disease; psoriasis; eczema; allergic conteat dermatitis; latex dermatitis; inflammatory bowel disease; ulcerative colitis; Crohn's disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 25-SEP-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (KRIE/) KRIEG A M. (BERG/) BERG D J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Immunostimulatory
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Berg DJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     administering an immunostimulatory nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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11-MAR-1998;
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11-MAR-1998;
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wound healing; obesity; diabetes; hearing loss;
cardiac insufficiency disorder; kidney disorder
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98US-0081105P
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disorder; kidney disorder; nervous system disorder;
disorder; kidney disorder; nervous system disorder;
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Pred. No. 9
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Ferrara N, 11
Goddard A, (13
Kljavin IJ,
Stewart TA,
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12-MAR-2000
130-MAY-2000
10-DEC-2000
24-AUG-2000
25-MAY-2000
26-FEB-2000
21-DEC-2000
21-DE
                                                                                                                                                                                                          The invention describes an isolated secreted and transmembrane (PRO) polypeptide (I). PRO337 polypeptide is useful for detecting PRO4993 polypeptide in a sample, and vice versa. PRO725, PRO700 and PRO739 are useful for detecting PRO1559 polypeptide in a sample, and PRO359 is useful for linking a bioactive molecule to a cell expressing a PRO337 is useful for linking a bioactive molecule to a cell expressing a PRO337 is useful for linking a bioactive molecule to a cell expressing a PRO4993 polypeptide. PRO1559 is useful for linking a bioactive molecule to a cell expressing a PRO4993 polypeptide. PRO1559 is useful for linking a bioactive molecule to a cell expressing a PRO4993 polypeptide, PRO35, PRO700 and PRO739 polypeptide, and PRO735, PRO700 and PRO739
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           isolated PRO polypeptides for example extracellular, secreted and brane bound proteins, useful for modulating the biological activities cells and for treating, for example diabetes, cancer, rheumatoid hritis, and hearing loss.
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